SEQUENCE LISTING

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<120> Transgenic plants carrying neoxanthin
      cleavege enzyme gene

<130> R3-102DP1

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<150> JP   2000-010056
<151> 2000-01-13

<160> 32

<170> PatentIn Ver. 2.0

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Met Val Ser Leu Leu Thr Met Pro Met Ser Gly Gly Ile Lys Thr Trp
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cct caa gcc caa att gat ttg ggt ttt agg ccc att aaa aga caa ccg   96
Pro Gln Ala Gln Ile Asp Leu Gly Phe Arg Pro Ile Lys Arg Gln Pro
            20   25   30

aag gtt att aaa tgc acg gtg cag atc gac gta acg gaa tta acc aaa   144
Lys Val Ile Lys Cys Thr Val Gln Ile Asp Val Thr Glu Leu Thr Lys
         35   40   45

aaa cgc caa tta ttt aca ccc aga acc acc gct act ccg ccg cag cat   192
Lys Arg Gln Leu Phe Thr Pro Arg Thr Thr Ala Thr Pro Pro Gln His
   50   55   60

aat cct ctc cgg cta aac atc ttc cag aaa gcg gcg gcg att gcg atc   240
Asn Pro Leu Arg Leu Asn Ile Phe Gln Lys Ala Ala Ala Ile Ala Ile
   65   70   75   80

gac gcg gct gag cgt gca tta atc tca cac gag caa gat tct cca ctt   288
Asp Ala Ala Glu Arg Ala Leu Ile Ser His Glu Gln Asp Ser Pro Leu
               85   90   95

ccc aaa acc gct gat cca cgt gtt cag att gcc ggg aat tat tcc ccg   336
Pro Lys Thr Ala Asp Pro Arg Val Gln Ile Ala Gly Asn Tyr Ser Pro
           100   105   110

gta ccg gaa tct tcc gtc cgg cga aac ctc acc gtc gaa gga aca atc   384
Val Pro Glu Ser Ser Val Arg Arg Asn Leu Thr Val Glu Gly Thr Ile
           115   120   125

cct gac tgc att gac ggt gtt tat atc cgt aac ggc gcg aat ccg atg   432
Pro Asp Cys Ile Asp Gly Val Tyr Ile Arg Asn Gly Ala Asn Pro Met
   130   135   140

ttt gag cca aca gct ggg cac cat tta ttc gac gga gac gga atg gtt   480
Phe Glu Pro Thr Ala Gly His His Leu Phe Asp Gly Asp Gly Met Val
145   150   155   160

cac gca gtt aaa ata acc aac ggt tca gct agc tac gca tgc cgg ttt   528
His Ala Val Lys Ile Thr Asn Gly Ser Ala Ser Tyr Ala Cys Arg Phe
           165   170   175

aca aaa acc gag aga ttg gtt cag gaa aaa cga ttg ggt cga cca gtt   576
Thr Lys Thr Glu Arg Leu Val Gln Glu Lys Arg Leu Gly Arg Pro Val
           180   185   190

ttc ccg aaa gca atc ggc gag ctt cac ggt cac tcg gga atc gca cgt   624
Phe Pro Lys Ala Ile Gly Glu Leu His Gly His Ser Gly Ile Ala Arg
          195   200   205

ttg atg ctg ttt tac gca cgt ggg ctt tgt ggt ctg atc aac aac caa   672
Leu Met Leu Phe Tyr Ala Arg Gly Leu Cys Gly Leu Ile Asn Asn Gln
    210   215   220

aac ggc gtc gga gta gca aac gcc ggt ttg gtt tac ttt aat aac cgg   720
Asn Gly Val Gly Val Ala Asn Ala Gly Leu Val Tyr Phe Asn Asn Arg
225   230   235   240

ctt tta gct atg tca gaa gac gat tta ccg tac caa tta aaa att act   768
Leu Leu Ala Met Ser Glu Asp Asp Leu Pro Tyr Gln Leu Lys Ile Thr
            245   250   255

caa acc ggc gat ctc caa acc gtt gga cgt tac gat ttc gac ggt cag   816
Gln Thr Gly Asp Leu Gln Thr Val Gly Arg Tyr Asp Phe Asp Gly Gln
            260   265   270

tta aaa tcc gca atg ata gct cac ccg aaa ctg gac ccg gtt acg aag   864
Leu Lys Ser Ala Met Ile Ala His Pro Lys Leu Asp Pro Val Thr Lys
            275   280   285

gag ctt cac gcg tta agc tac gac gtc gtt aag aaa cct tac ctg aaa   912
Glu Leu His Ala Leu Ser Tyr Asp Val Val Lys Lys Pro Tyr Leu Lys
    290   295   300

tac ttc aga ttc tcg cca gac ggc gtt aaa tcg ccg gaa ttg gag atc   960
Tyr Phe Arg Phe Ser Pro Asp Gly Val Lys Ser Pro Glu Leu Glu Ile
305   310   315   320

ccg ctc gaa act ccg acg atg att cac gat ttc gct ata acg gag aat   1008
Pro Leu Glu Thr Pro Thr Met Ile His Asp Phe Ala Ile Thr Glu Asn
                325   330   335

ttt gtg gtg att cct gat caa caa gtc gtg ttc aag ctc ggc gag atg   1056
Phe Val Val Ile Pro Asp Gln Gln Val Val Phe Lys Leu Gly Glu Met
            340   345   350

att tcc ggt aaa tct ccg gtt gtt ttc gac gga gaa aag gtt tcc cga   1104
Ile Ser Gly Lys Ser Pro Val Val Phe Asp Gly Glu Lys Val Ser Arg
    355   360   365

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ttg ggg ata atg ccc aag gac gcg aca gaa gct tct cag ata atc tgg    1152
Leu Gly Ile Met Pro Lys Asp Ala Thr Glu Ala Ser Gln Ile Ile Trp
    370             375             380

gtg aac tct ccg gag acg ttc tgt ttt cat ctc tgg aat gca tgg gaa    1200
Val Asn Ser Pro Glu Thr Phe Cys Phe His Leu Trp Asn Ala Trp Glu
385             390             395             400

tcg ccg gag acg gag gag att gtg gtg atc gga tcg tgt atg tcg ccg    1248
Ser Pro Glu Thr Glu Glu Ile Val Val Ile Gly Ser Cys Met Ser Pro
            405             410             415

gcg gat tca atc ttc aac gag aga gac gag agc ttg aga agc gtt ttg    1296
Ala Asp Ser Ile Phe Asn Glu Arg Asp Glu Ser Leu Arg Ser Val Leu
            420             425             430

tcg gag atc agg ata aac ctc aga aca cgt aaa acc acg cgt cgt tcg    1344
Ser Glu Ile Arg Ile Asn Leu Arg Thr Arg Lys Thr Thr Arg Arg Ser
            435             440             445

ttg ttg gtt aac gag gat gta aat tta gag att ggt atg gtt aac cgg    1392
Leu Leu Val Asn Glu Asp Val Asn Leu Glu Ile Gly Met Val Asn Arg
    450             455             460

aac cgg tta gga aga aaa acc cgg ttc gcg ttt ttg gct att gct tat    1440
Asn Arg Leu Gly Arg Lys Thr Arg Phe Ala Phe Leu Ala Ile Ala Tyr
465             470             475             480

cct tgg cca aaa gtt tcc ggt ttc gct aag gtc gat ctt tgc acc ggt    1488
Pro Trp Pro Lys Val Ser Gly Phe Ala Lys Val Asp Leu Cys Thr Gly
            485             490             495

gag atg aaa aaa tat att tac ggc ggt gag aaa tat ggc ggc gaa ccg    1536
Glu Met Lys Lys Tyr Ile Tyr Gly Gly Glu Lys Tyr Gly Gly Glu Pro
            500             505             510

ttt ttc ttg ccc ggc aac tcc ggt aac ggc gaa gaa aat gaa gat gac    1584
Phe Phe Leu Pro Gly Asn Ser Gly Asn Gly Glu Glu Asn Glu Asp Asp
            515             520             525
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ggt tat ata ttt tgt cac gtt cat gac gaa gaa aca aag aca tca gag    1632
Gly Tyr Ile Phe Cys His Val His Asp Glu Glu Thr Lys Thr Ser Glu
    530             535             540

ctt cag att att aac gct gtt aat tta aag ctt gaa gct acg att aaa    1680
Leu Gln Ile Ile Asn Ala Val Asn Leu Lys Leu Glu Ala Thr Ile Lys
545             550             555             560

cta ccg tct aga gta ccg tat ggg ttt cat ggc aca ttt gtg gat tcg    1728
Leu Pro Ser Arg Val Pro Tyr Gly Phe His Gly Thr Phe Val Asp Ser
            565             570             575

aat gaa ctc gtt gat caa tta taa                                    1752
Asn Glu Leu Val Asp Gln Leu
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Pro Gln Ala Gln Ile Asp Leu Gly Phe Arg Pro Ile Lys Arg Gln Pro
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Lys Val Ile Lys Cys Thr Val Gln Ile Asp Val Thr Glu Leu Thr Lys
        35              40              45

Lys Arg Gln Leu Phe Thr Pro Arg Thr Thr Ala Thr Pro Pro Gln His
        50              55              60

Asn Pro Leu Arg Leu Asn Ile Phe Gln Lys Ala Ala Ala Ile Ala Ile
65              70              75              80

Asp Ala Ala Glu Arg Ala Leu Ile Ser His Glu Gln Asp Ser Pro Leu
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                        85   90   95

Pro Lys Thr Ala Asp Pro Arg Val Gln Ile Ala Gly Asn Tyr Ser Pro
          100   105   110

Val Pro Glu Ser Ser Val Arg Arg Asn Leu Thr Val Glu Gly Thr Ile
          115   120   125

Pro Asp Cys Ile Asp Gly Val Tyr Ile Arg Asn Gly Ala Asn Pro Met
          130   135   140

Phe Glu Pro Thr Ala Gly His His Leu Phe Asp Gly Asp Gly Met Val
145   150   155   160

His Ala Val Lys Ile Thr Asn Gly Ser Ala Ser Tyr Ala Cys Arg Phe
              165   170   175

Thr Lys Thr Glu Arg Leu Val Gln Glu Lys Arg Leu Gly Arg Pro Val
              180   185   190

Phe Pro Lys Ala Ile Gly Glu Leu His Gly His Ser Gly Ile Ala Arg
          195   200   205

Leu Met Leu Phe Tyr Ala Arg Gly Leu Cys Gly Leu Ile Asn Asn Gln
          210   215   220

Asn Gly Val Gly Val Ala Asn Ala Gly Leu Val Tyr Phe Asn Asn Arg
225   230   235   240

Leu Leu Ala Met Ser Glu Asp Asp Leu Pro Tyr Gln Leu Lys Ile Thr
              245   250   255

Gln Thr Gly Asp Leu Gln Thr Val Gly Arg Tyr Asp Phe Asp Gly Gln
          260   265   270

Leu Lys Ser Ala Met Ile Ala His Pro Lys Leu Asp Pro Val Thr Lys
          275   280   285

Glu Leu His Ala Leu Ser Tyr Asp Val Val Lys Lys Pro Tyr Leu Lys
          290   295   300

Tyr Phe Arg Phe Ser Pro Asp Gly Val Lys Ser Pro Glu Leu Glu Ile
305   310   315   320

Pro Leu Glu Thr Pro Thr Met Ile His Asp Phe Ala Ile Thr Glu Asn
                325   330   335

Phe Val Val Ile Pro Asp Gln Gln Val Val Phe Lys Leu Gly Glu Met
            340   345   350

Ile Ser Gly Lys Ser Pro Val Val Phe Asp Gly Glu Lys Val Ser Arg
            355   360   365

Leu Gly Ile Met Pro Lys Asp Ala Thr Glu Ala Ser Gln Ile Ile Trp
        370   375   380

Val Asn Ser Pro Glu Thr Phe Cys Phe His Leu Trp Asn Ala Trp Glu
385   390   395   400

Ser Pro Glu Thr Glu Glu Ile Val Val Ile Gly Ser Cys Met Ser Pro
                405   410   415

Ala Asp Ser Ile Phe Asn Glu Arg Asp Glu Ser Leu Arg Ser Val Leu
            420   425   430

Ser Glu Ile Arg Ile Asn Leu Arg Thr Arg Lys Thr Thr Arg Arg Ser
        435   440   445

Leu Leu Val Asn Glu Asp Val Asn Leu Glu Ile Gly Met Val Asn Arg
        450   455   460

Asn Arg Leu Gly Arg Lys Thr Arg Phe Ala Phe Leu Ala Ile Ala Tyr
465   470   475   480

Pro Trp Pro Lys Val Ser Gly Phe Ala Lys Val Asp Leu Cys Thr Gly
            485   490   495

Glu Met Lys Lys Tyr Ile Tyr Gly Gly Glu Lys Tyr Gly Gly Glu Pro
            500   505   510

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Phe Phe Leu Pro Gly Asn Ser Gly Asn Gly Glu Glu Asn Glu Asp Asp
        515             520             525

Gly Tyr Ile Phe Cys His Val His Asp Glu Glu Thr Lys Thr Ser Glu
    530             535             540

Leu Gln Ile Ile Asn Ala Val Asn Leu Lys Leu Glu Ala Thr Ile Lys
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Leu Pro Ser Arg Val Pro Tyr Gly Phe His Gly Thr Phe Val Asp Ser
            565             570             575

Asn Glu Leu Val Asp Gln Leu
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Met Asp Ser Val Ser Ser Ser Ser Phe Leu Ser Ser Thr Phe Ser Leu
  1               5              10              15

cat cac tct ctt ctt cgc cgc cga tct tcc tct cct act ctc ctc cgt    96
His His Ser Leu Leu Arg Arg Arg Ser Ser Ser Pro Thr Leu Leu Arg
            20              25              30

atc aac tcc gcc gtc gtc gaa gaa cgt tct cca atc aca aac cca agc   144
Ile Asn Ser Ala Val Val Glu Glu Arg Ser Pro Ile Thr Asn Pro Ser
        35              40              45

gac aac aat gat cgt cgt aac aaa ccc aaa aca ctc cac aac cga acc   192
Asp Asn Asn Asp Arg Arg Asn Lys Pro Lys Thr Leu His Asn Arg Thr
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                50                      55                      60

aat cac acc tta gtc tca tca cca ccg aaa ctc cga cca gaa atg act     240
Asn His Thr Leu Val Ser Ser Pro Pro Lys Leu Arg Pro Glu Met Thr
 65                      70                      75                80

ctc gca aca gct ctc ttc acc acc gtc gaa gat gta atc aac acg ttc     288
Leu Ala Thr Ala Leu Phe Thr Thr Val Glu Asp Val Ile Asn Thr Phe
                     85                      90                      95

atc gat cca cct tca cgt cct tcc gtt gat cca aaa cat gtc ctc tct     336
Ile Asp Pro Pro Ser Arg Pro Ser Val Asp Pro Lys His Val Leu Ser
                    100                     105                     110

gat aac ttc gct cct gtc ctc gac gag ctt cct cca aca gac tgt gaa     384
Asp Asn Phe Ala Pro Val Leu Asp Glu Leu Pro Pro Thr Asp Cys Glu
                    115                     120                     125

atc atc cac ggc act ctt cca ctg tca ctt aac ggc gct tac atc cgt     432
Ile Ile His Gly Thr Leu Pro Leu Ser Leu Asn Gly Ala Tyr Ile Arg
               130                     135                     140

aac ggt cca aat cca cag ttt ctc cct cgt ggt cct tac cat ctc ttc     480
Asn Gly Pro Asn Pro Gln Phe Leu Pro Arg Gly Pro Tyr His Leu Phe
145                     150                     155                 160

gac ggc gac ggt atg ctt cac gcc ata aaa atc cac aac ggt aaa gcc     528
Asp Gly Asp Gly Met Leu His Ala Ile Lys Ile His Asn Gly Lys Ala
                    165                     170                     175

act ctc tgt agc aga tac gtc aag act tat aaa tac aac gtc gag aaa     576
Thr Leu Cys Ser Arg Tyr Val Lys Thr Tyr Lys Tyr Asn Val Glu Lys
                    180                     185                     190

caa acc gga gct ccg gtt atg cct aac gtg ttt tcc gga ttc aac ggt     624
Gln Thr Gly Ala Pro Val Met Pro Asn Val Phe Ser Gly Phe Asn Gly
               195                     200                     205

gta acg gcg tca gta gct cgt gga gct tta acg gca gct agg gtt tta     672
Val Thr Ala Ser Val Ala Arg Gly Ala Leu Thr Ala Ala Arg Val Leu
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| | 210 | | 215 | | 220 | |

acc gga cag tat aat ccg gtt aac ggc att ggt tta gct aat aca agt   720
Thr Gly Gln Tyr Asn Pro Val Asn Gly Ile Gly Leu Ala Asn Thr Ser
225   230   235   240

cta gct ttc ttc agt aac cgt ctc ttt gct tta ggt gaa tct gat tta   768
Leu Ala Phe Phe Ser Asn Arg Leu Phe Ala Leu Gly Glu Ser Asp Leu
                245   250   255

ccc tac gcc gtc cga tta acc gaa tca gga gat att gaa acg atc gga   816
Pro Tyr Ala Val Arg Leu Thr Glu Ser Gly Asp Ile Glu Thr Ile Gly
                260   265   270

cgg tac gat ttc gac ggg aaa tta gcg atg agt atg aca gct cat cct   864
Arg Tyr Asp Phe Asp Gly Lys Leu Ala Met Ser Met Thr Ala His Pro
                275   280   285

aaa acc gat cca ata acc gga gaa act ttc gct ttc cgg tac ggt ccg   912
Lys Thr Asp Pro Ile Thr Gly Glu Thr Phe Ala Phe Arg Tyr Gly Pro
            290   295   300

gtt cca ccg ttt tta aca tat ttc cgg ttt gat tcc gcc ggg aaa aaa   960
Val Pro Pro Phe Leu Thr Tyr Phe Arg Phe Asp Ser Ala Gly Lys Lys
305   310   315   320

caa aga gac gtt ccg ata ttc tcg atg acg tct ccg tcg ttt ctc cat   1008
Gln Arg Asp Val Pro Ile Phe Ser Met Thr Ser Pro Ser Phe Leu His
                325   330   335

gac ttc gcg atc acg aaa cgt cac gcg att ttc gca gag att cag ctt   1056
Asp Phe Ala Ile Thr Lys Arg His Ala Ile Phe Ala Glu Ile Gln Leu
                340   345   350

ggc atg agg atg aac atg ttg gat ttg gtt ctc gaa ggt ggt tct ccg   1104
Gly Met Arg Met Asn Met Leu Asp Leu Val Leu Glu Gly Gly Ser Pro
                355   360   365

gtt ggt act gat aac gga aaa act cca agg ctt gga gtg att cct aag   1152
Val Gly Thr Asp Asn Gly Lys Thr Pro Arg Leu Gly Val Ile Pro Lys

370   375   380

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tac gcc gga gat gag tcg gag atg aaa tgg ttc gaa gtt cct gga ttc     1200
Tyr Ala Gly Asp Glu Ser Glu Met Lys Trp Phe Glu Val Pro Gly Phe
385             390             395             400
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aat atc att cac gct att aat gct tgg gat gaa gat gat gga aac agc     1248
Asn Ile Ile His Ala Ile Asn Ala Trp Asp Glu Asp Asp Gly Asn Ser
            405             410             415
```

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gtc gtt ttg att gca ccg aat att atg tcg att gaa cat act tta gag     1296
Val Val Leu Ile Ala Pro Asn Ile Met Ser Ile Glu His Thr Leu Glu
            420             425             430
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```
agg atg gat ctg gtt cat gct ttg gtg gag aag gtg aag atc gat ctc     1344
Arg Met Asp Leu Val His Ala Leu Val Glu Lys Val Lys Ile Asp Leu
            435             440             445
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gtc acc ggg att gtg aga cgt cat ccg atc tca gcg agg aat ctc gat     1392
Val Thr Gly Ile Val Arg Arg His Pro Ile Ser Ala Arg Asn Leu Asp
    450             455             460
```

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ttc gct gtg att aat ccg gcg ttt ctc ggg aga tgt agc agg tac gtt     1440
Phe Ala Val Ile Asn Pro Ala Phe Leu Gly Arg Cys Ser Arg Tyr Val
465             470             475             480
```

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tac gcg gcg att gga gat ccg atg ccg aag atc tcc ggt gtg gtg aag     1488
Tyr Ala Ala Ile Gly Asp Pro Met Pro Lys Ile Ser Gly Val Val Lys
            485             490             495
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ctt gat gtg tct aaa gga gat cgg gat gat tgt acg gtg gcc cgt aga     1536
Leu Asp Val Ser Lys Gly Asp Arg Asp Asp Cys Thr Val Ala Arg Arg
            500             505             510
```

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atg tac ggt tca ggt tgt tac ggc gga gaa ccg ttt ttc gta gct agg     1584
Met Tyr Gly Ser Gly Cys Tyr Gly Gly Glu Pro Phe Phe Val Ala Arg
            515             520             525
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gat cct ggt aat ccg gag gcg gag gag gat gat ggt tat gtg gtg acg     1632
Asp Pro Gly Asn Pro Glu Ala Glu Glu Asp Asp Gly Tyr Val Val Thr
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```
          530                    535                    540

tat gtt cac gat gaa gtg act gga gaa tcg aag ttt ctg gtg atg gac    1680
Tyr Val His Asp Glu Val Thr Gly Glu Ser Lys Phe Leu Val Met Asp
545                550                555                560

gct aaa tcg ccg gag ctt gaa atc gtc gcc gcc gtg agg ttg ccg cga    1728
Ala Lys Ser Pro Glu Leu Glu Ile Val Ala Ala Val Arg Leu Pro Arg
               565                570                575

agg gtt ccg tac gga ttc cat ggg tta ttt gtc aag gaa agt gac ctt    1776
Arg Val Pro Tyr Gly Phe His Gly Leu Phe Val Lys Glu Ser Asp Leu
               580                585                590

aat aag ctt taa                                                    1788
Asn Lys Leu
          595
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             20                 25                 30

Ile Asn Ser Ala Val Val Glu Glu Arg Ser Pro Ile Thr Asn Pro Ser
             35                 40                 45

Asp Asn Asn Asp Arg Arg Asn Lys Pro Lys Thr Leu His Asn Arg Thr
        50                 55                 60

Asn His Thr Leu Val Ser Ser Pro Pro Lys Leu Arg Pro Glu Met Thr
 65                 70                 75                 80
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Leu Ala Thr Ala Leu Phe Thr Thr Val Glu Asp Val Ile Asn Thr Phe
                85                  90                  95

Ile Asp Pro Pro Ser Arg Pro Ser Val Asp Pro Lys His Val Leu Ser
               100                 105                 110

Asp Asn Phe Ala Pro Val Leu Asp Glu Leu Pro Pro Thr Asp Cys Glu
            115                 120                 125

Ile Ile His Gly Thr Leu Pro Leu Ser Leu Asn Gly Ala Tyr Ile Arg
        130                 135                 140

Asn Gly Pro Asn Pro Gln Phe Leu Pro Arg Gly Pro Tyr His Leu Phe
145                 150                 155                 160

Asp Gly Asp Gly Met Leu His Ala Ile Lys Ile His Asn Gly Lys Ala
                165                 170                 175

Thr Leu Cys Ser Arg Tyr Val Lys Thr Tyr Lys Tyr Asn Val Glu Lys
            180                 185                 190

Gln Thr Gly Ala Pro Val Met Pro Asn Val Phe Ser Gly Phe Asn Gly
            195                 200                 205

Val Thr Ala Ser Val Ala Arg Gly Ala Leu Thr Ala Ala Arg Val Leu
    210                 215                 220

Thr Gly Gln Tyr Asn Pro Val Asn Gly Ile Gly Leu Ala Asn Thr Ser
225                 230                 235                 240

Leu Ala Phe Phe Ser Asn Arg Leu Phe Ala Leu Gly Glu Ser Asp Leu
                245                 250                 255

Pro Tyr Ala Val Arg Leu Thr Glu Ser Gly Asp Ile Glu Thr Ile Gly
                260                 265                 270

Arg Tyr Asp Phe Asp Gly Lys Leu Ala Met Ser Met Thr Ala His Pro
            275                 280                 285

Lys Thr Asp Pro Ile Thr Gly Glu Thr Phe Ala Phe Arg Tyr Gly Pro
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                290                    295                    300

Val Pro Pro Phe Leu Thr Tyr Phe Arg Phe Asp Ser Ala Gly Lys Lys
305                310                315                320

Gln Arg Asp Val Pro Ile Phe Ser Met Thr Ser Pro Ser Phe Leu His
                325                330                335

Asp Phe Ala Ile Thr Lys Arg His Ala Ile Phe Ala Glu Ile Gln Leu
                340                345                350

Gly Met Arg Met Asn Met Leu Asp Leu Val Leu Glu Gly Gly Ser Pro
                355                360                365

Val Gly Thr Asp Asn Gly Lys Thr Pro Arg Leu Gly Val Ile Pro Lys
    370                375                380

Tyr Ala Gly Asp Glu Ser Glu Met Lys Trp Phe Glu Val Pro Gly Phe
385                390                395                400

Asn Ile Ile His Ala Ile Asn Ala Trp Asp Glu Asp Asp Gly Asn Ser
                405                410                415

Val Val Leu Ile Ala Pro Asn Ile Met Ser Ile Glu His Thr Leu Glu
                420                425                430

Arg Met Asp Leu Val His Ala Leu Val Glu Lys Val Lys Ile Asp Leu
                435                440                445

Val Thr Gly Ile Val Arg Arg His Pro Ile Ser Ala Arg Asn Leu Asp
    450                455                460

Phe Ala Val Ile Asn Pro Ala Phe Leu Gly Arg Cys Ser Arg Tyr Val
465                470                475                480

Tyr Ala Ala Ile Gly Asp Pro Met Pro Lys Ile Ser Gly Val Val Lys
                485                490                495

Leu Asp Val Ser Lys Gly Asp Arg Asp Asp Cys Thr Val Ala Arg Arg
                500                505                510
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Met Tyr Gly Ser Gly Cys Tyr Gly Gly Glu Pro Phe Phe Val Ala Arg
        515             520             525

Asp Pro Gly Asn Pro Glu Ala Glu Glu Asp Asp Gly Tyr Val Val Thr
    530             535             540

Tyr Val His Asp Glu Val Thr Gly Glu Ser Lys Phe Leu Val Met Asp
545             550             555             560

Ala Lys Ser Pro Glu Leu Glu Ile Val Ala Ala Val Arg Leu Pro Arg
            565             570             575

Arg Val Pro Tyr Gly Phe His Gly Leu Phe Val Lys Glu Ser Asp Leu
        580             585             590

Asn Lys Leu
        595
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Met Ala Ser Phe Thr Ala Thr Ala Ala Val Ser Gly Arg Trp Leu Gly
 1               5                  10                  15

ggc aat cat act cag ccg cca tta tcg tct tct caa agc tcc gac ttg    96
Gly Asn His Thr Gln Pro Pro Leu Ser Ser Ser Gln Ser Ser Asp Leu
            20                  25                  30

agt tat tgt agc tcc tta cct atg gcc agt cgt gtc aca cgt aag ctc   144
Ser Tyr Cys Ser Ser Leu Pro Met Ala Ser Arg Val Thr Arg Lys Leu
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                35                      40                      45

aat gtt tca tct gcg ctt cac act cct cca gct ctt cat ttc cct aag    192
Asn Val Ser Ser Ala Leu His Thr Pro Pro Ala Leu His Phe Pro Lys
        50                      55                      60


caa tca tca aac tct ccc gcc att gtt gtt aag ccc aaa gcc aaa gaa    240
Gln Ser Ser Asn Ser Pro Ala Ile Val Val Lys Pro Lys Ala Lys Glu
 65                      70                      75                 80


tcc aac act aaa cag atg aat ttg ttc cag aga gcg gcg gcg gca gcg    288
Ser Asn Thr Lys Gln Met Asn Leu Phe Gln Arg Ala Ala Ala Ala Ala
                85                      90                      95


ttg gac gcg gcg gag ggt ttc ctt gtc agc cac gag aag cta cac ccg    336
Leu Asp Ala Ala Glu Gly Phe Leu Val Ser His Glu Lys Leu His Pro
            100                     105                     110


ctt cct aaa acg gct gat cct agt gtt cag atc gcc gga aat ttt gct    384
Leu Pro Lys Thr Ala Asp Pro Ser Val Gln Ile Ala Gly Asn Phe Ala
            115                     120                     125


ccg gtg aat gaa cag ccc gtc cgg cgt aat ctt ccg gtg gtc gga aaa    432
Pro Val Asn Glu Gln Pro Val Arg Arg Asn Leu Pro Val Val Gly Lys
        130                     135                     140


ctt ccc gat tcc atc aaa gga gtg tat gtg cgc aac gga gct aac cca    480
Leu Pro Asp Ser Ile Lys Gly Val Tyr Val Arg Asn Gly Ala Asn Pro
145                     150                     155                 160


ctt cac gag ccg gtg aca ggt cac cac ttc ttc gac gga gac ggt atg    528
Leu His Glu Pro Val Thr Gly His His Phe Phe Asp Gly Asp Gly Met
            165                     170                     175


gtt cac gcc gtc aaa ttc gaa cac ggt tca gct agc tac gct tgc cgg    576
Val His Ala Val Lys Phe Glu His Gly Ser Ala Ser Tyr Ala Cys Arg
            180                     185                     190


ttt act cag act aac cgg ttt gtt cag gaa cgt caa ttg ggt cga ccg    624
Phe Thr Gln Thr Asn Arg Phe Val Gln Glu Arg Gln Leu Gly Arg Pro
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            195                    200                    205

gtt ttc ccc aaa gcc atc ggt gag ctt cac ggc cac acc ggt att gcc     672
Val Phe Pro Lys Ala Ile Gly Glu Leu His Gly His Thr Gly Ile Ala
    210                    215                    220


cga ctc atg cta ttc tac gcc aga gct gca gcc ggt ata gtc gac ccg     720
Arg Leu Met Leu Phe Tyr Ala Arg Ala Ala Ala Gly Ile Val Asp Pro
225                    230                    235                    240


gca cac gga acc ggt gta gct aac gcc ggt ttg gtc tat ttc aat ggc     768
Ala His Gly Thr Gly Val Ala Asn Ala Gly Leu Val Tyr Phe Asn Gly
                   245                    250                    255


cgg tta ttg gct atg tcg gag gat gat tta cct tac caa gtt cag atc     816
Arg Leu Leu Ala Met Ser Glu Asp Asp Leu Pro Tyr Gln Val Gln Ile
                   260                    265                    270


act ccc aat gga gat tta aaa acc gtt ggt cgg ttc gat ttt gat gga     864
Thr Pro Asn Gly Asp Leu Lys Thr Val Gly Arg Phe Asp Phe Asp Gly
                   275                    280                    285


caa tta gaa tcc aca atg att gcc cac ccg aaa gtc gac ccg gaa tcc     912
Gln Leu Glu Ser Thr Met Ile Ala His Pro Lys Val Asp Pro Glu Ser
    290                    295                    300


ggt gaa ctc ttc gct tta agc tac gac gtc gtt tca aag cct tac cta     960
Gly Glu Leu Phe Ala Leu Ser Tyr Asp Val Val Ser Lys Pro Tyr Leu
305                    310                    315                    320


aaa tac ttc cga ttc tca ccg gac gga act aaa tca ccg gac gtc gag    1008
Lys Tyr Phe Arg Phe Ser Pro Asp Gly Thr Lys Ser Pro Asp Val Glu
                   325                    330                    335


att cag ctt gat cag cca acg atg atg cac gat ttc gcg att aca gag    1056
Ile Gln Leu Asp Gln Pro Thr Met Met His Asp Phe Ala Ile Thr Glu
                   340                    345                    350


aac ttc gtc gtc gta cct gac cag caa gtc gtt ttc aag ctg ccg gag    1104
Asn Phe Val Val Val Pro Asp Gln Gln Val Val Phe Lys Leu Pro Glu
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355   360   365

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atg atc cgc ggt ggg tct ccg gtg gtt tac gac aag aac aag gtc gca     1152
Met Ile Arg Gly Gly Ser Pro Val Val Tyr Asp Lys Asn Lys Val Ala
    370             375             380

aga ttc ggg att tta gac aaa tac gcc gaa gat tca tcg aac att aag     1200
Arg Phe Gly Ile Leu Asp Lys Tyr Ala Glu Asp Ser Ser Asn Ile Lys
385             390             395             400

tgg att gat gct cca gat tgc ttc tgc ttc cat ctc tgg aac gct tgg     1248
Trp Ile Asp Ala Pro Asp Cys Phe Cys Phe His Leu Trp Asn Ala Trp
            405             410             415

gaa gag cca gaa aca gat gaa gtc gtc gtg ata ggg tcc tgt atg act     1296
Glu Glu Pro Glu Thr Asp Glu Val Val Val Ile Gly Ser Cys Met Thr
            420             425             430

cca cca gac tca att ttc aac gag tct gac gag aat ctc aag agt gtc     1344
Pro Pro Asp Ser Ile Phe Asn Glu Ser Asp Glu Asn Leu Lys Ser Val
            435             440             445

ctg tct gaa atc cgc ctg aat ctc aaa acc ggt gaa tca act cgc cgt     1392
Leu Ser Glu Ile Arg Leu Asn Leu Lys Thr Gly Glu Ser Thr Arg Arg
    450             455             460

ccg atc atc tcc aac gaa gat caa caa gtc aac ctc gaa gca ggg atg     1440
Pro Ile Ile Ser Asn Glu Asp Gln Gln Val Asn Leu Glu Ala Gly Met
465             470             475             480

gtc aac aga aac atg ctc ggc cgt aaa acc aaa ttc gct tac ttg gct     1488
Val Asn Arg Asn Met Leu Gly Arg Lys Thr Lys Phe Ala Tyr Leu Ala
            485             490             495

tta gcc gag ccg tgg cct aaa gtc tca gga ttc gct aaa gtt gat ctc     1536
Leu Ala Glu Pro Trp Pro Lys Val Ser Gly Phe Ala Lys Val Asp Leu
            500             505             510

act act gga gaa gtt aag aaa cat ctt tac ggc gat aac cgt tac gga     1584
Thr Thr Gly Glu Val Lys Lys His Leu Tyr Gly Asp Asn Arg Tyr Gly
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                515                520                525

gga gag cct ctg ttt ctc ccc gga gaa gga gga gag gaa gac gaa gga    1632
Gly Glu Pro Leu Phe Leu Pro Gly Glu Gly Gly Glu Glu Asp Glu Gly
    530                535                540

tac atc ctc tgt ttc gtt cac gac gag aag aca tgg aaa tcg gag tta    1680
Tyr Ile Leu Cys Phe Val His Asp Glu Lys Thr Trp Lys Ser Glu Leu
545                550                555                560

cag ata gtt aac gcc gtt agc tta gag gtt gaa gca acg gtt aaa ctt    1728
Gln Ile Val Asn Ala Val Ser Leu Glu Val Glu Ala Thr Val Lys Leu
                565                570                575

ccg tca agg gtt ccg tac gga ttt cac ggt aca ttc atc gga gcc gat    1776
Pro Ser Arg Val Pro Tyr Gly Phe His Gly Thr Phe Ile Gly Ala Asp
                580                585                590

gat ttg gcg aag cag gtc gtg tga                                    1800
Asp Leu Ala Lys Gln Val Val
                595                600
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<211> 599
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<213> Arabidopsis thaliana

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  1                5                10                15

Gly Asn His Thr Gln Pro Pro Leu Ser Ser Ser Gln Ser Ser Asp Leu
            20                25                30

Ser Tyr Cys Ser Ser Leu Pro Met Ala Ser Arg Val Thr Arg Lys Leu
        35                40                45

Asn Val Ser Ser Ala Leu His Thr Pro Pro Ala Leu His Phe Pro Lys
    50                55                60
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Gln Ser Ser Asn Ser Pro Ala Ile Val Val Lys Pro Lys Ala Lys Glu
65   70   75   80

Ser Asn Thr Lys Gln Met Asn Leu Phe Gln Arg Ala Ala Ala Ala Ala
              85   90   95

Leu Asp Ala Ala Glu Gly Phe Leu Val Ser His Glu Lys Leu His Pro
            100   105   110

Leu Pro Lys Thr Ala Asp Pro Ser Val Gln Ile Ala Gly Asn Phe Ala
            115   120   125

Pro Val Asn Glu Gln Pro Val Arg Arg Asn Leu Pro Val Val Gly Lys
          130   135   140

Leu Pro Asp Ser Ile Lys Gly Val Tyr Val Arg Asn Gly Ala Asn Pro
145   150   155   160

Leu His Glu Pro Val Thr Gly His His Phe Phe Asp Gly Asp Gly Met
              165   170   175

Val His Ala Val Lys Phe Glu His Gly Ser Ala Ser Tyr Ala Cys Arg
            180   185   190

Phe Thr Gln Thr Asn Arg Phe Val Gln Glu Arg Gln Leu Gly Arg Pro
          195   200   205

Val Phe Pro Lys Ala Ile Gly Glu Leu His Gly His Thr Gly Ile Ala
        210   215   220

Arg Leu Met Leu Phe Tyr Ala Arg Ala Ala Ala Gly Ile Val Asp Pro
225   230   235   240

Ala His Gly Thr Gly Val Ala Asn Ala Gly Leu Val Tyr Phe Asn Gly
              245   250   255

Arg Leu Leu Ala Met Ser Glu Asp Asp Leu Pro Tyr Gln Val Gln Ile
            260   265   270

Thr Pro Asn Gly Asp Leu Lys Thr Val Gly Arg Phe Asp Phe Asp Gly
    275   280   285

Gln Leu Glu Ser Thr Met Ile Ala His Pro Lys Val Asp Pro Glu Ser
    290   295   300

Gly Glu Leu Phe Ala Leu Ser Tyr Asp Val Val Ser Lys Pro Tyr Leu
305   310   315   320

Lys Tyr Phe Arg Phe Ser Pro Asp Gly Thr Lys Ser Pro Asp Val Glu
         325   330   335

Ile Gln Leu Asp Gln Pro Thr Met Met His Asp Phe Ala Ile Thr Glu
      340   345   350

Asn Phe Val Val Val Pro Asp Gln Gln Val Val Phe Lys Leu Pro Glu
      355   360   365

Met Ile Arg Gly Gly Ser Pro Val Val Tyr Asp Lys Asn Lys Val Ala
    370   375   380

Arg Phe Gly Ile Leu Asp Lys Tyr Ala Glu Asp Ser Ser Asn Ile Lys
385   390   395   400

Trp Ile Asp Ala Pro Asp Cys Phe Cys Phe His Leu Trp Asn Ala Trp
        405   410   415

Glu Glu Pro Glu Thr Asp Glu Val Val Val Ile Gly Ser Cys Met Thr
        420   425   430

Pro Pro Asp Ser Ile Phe Asn Glu Ser Asp Glu Asn Leu Lys Ser Val
      435   440   445

Leu Ser Glu Ile Arg Leu Asn Leu Lys Thr Gly Glu Ser Thr Arg Arg
    450   455   460

Pro Ile Ile Ser Asn Glu Asp Gln Gln Val Asn Leu Glu Ala Gly Met
465   470   475   480

Val Asn Arg Asn Met Leu Gly Arg Lys Thr Lys Phe Ala Tyr Leu Ala

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          485                 490                 495

Leu Ala Glu Pro Trp Pro Lys Val Ser Gly Phe Ala Lys Val Asp Leu
          500                 505                 510


Thr Thr Gly Glu Val Lys Lys His Leu Tyr Gly Asp Asn Arg Tyr Gly
          515                 520                 525


Gly Glu Pro Leu Phe Leu Pro Gly Glu Gly Gly Glu Glu Asp Glu Gly
          530                 535                 540


Tyr Ile Leu Cys Phe Val His Asp Glu Lys Thr Trp Lys Ser Glu Leu
545                 550                 555                 560


Gln Ile Val Asn Ala Val Ser Leu Glu Val Glu Ala Thr Val Lys Leu
              565                 570                 575


Pro Ser Arg Val Pro Tyr Gly Phe His Gly Thr Phe Ile Gly Ala Asp
              580                 585                 590


Asp Leu Ala Lys Gln Val Val
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Met Ala Glu Lys Leu Ser Asp Gly Ser Ser Ile Ile Ser Val His Pro
 1               5                  10                  15


aga ccc tcc aag ggt ttc tcc tcg aag ctt ctc gat ctt ctc gag aga    96
Arg Pro Ser Lys Gly Phe Ser Ser Lys Leu Leu Asp Leu Leu Glu Arg
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                    20   25   30

ctt gtt gtc aag ctc atg cac gat gct tct ctc cct ctc cac tac ctc   144
Leu Val Val Lys Leu Met His Asp Ala Ser Leu Pro Leu His Tyr Leu
            35   40   45

tca ggc aac ttc gct ccc atc cgt gat gaa act cct ccc gtc aag gat   192
Ser Gly Asn Phe Ala Pro Ile Arg Asp Glu Thr Pro Pro Val Lys Asp
        50   55   60

ctc ccc gtc cat gga ttt ctt ccc gaa tgc ttg aat ggt gaa ttt gtg   240
Leu Pro Val His Gly Phe Leu Pro Glu Cys Leu Asn Gly Glu Phe Val
65   70   75   80

agg gtt ggt cca aac ccc aag ttt gat gct gtc gct gga tat cac tgg   288
Arg Val Gly Pro Asn Pro Lys Phe Asp Ala Val Ala Gly Tyr His Trp
                85   90   95

ttt gat gga gat ggg atg att cat ggg gta cgc atc aaa gat ggg aaa   336
Phe Asp Gly Asp Gly Met Ile His Gly Val Arg Ile Lys Asp Gly Lys
            100   105   110

gct act tat gtt tct cga tat gtt aag aca tca cgt ctt aag cag gaa   384
Ala Thr Tyr Val Ser Arg Tyr Val Lys Thr Ser Arg Leu Lys Gln Glu
            115   120   125

gag ttc ttc gga gct gcc aaa ttc atg aag att ggt gac ctt aag ggg   432
Glu Phe Phe Gly Ala Ala Lys Phe Met Lys Ile Gly Asp Leu Lys Gly
        130   135   140

ttt ttc gga ttg cta atg gtc aat gtc caa cag ctg aga acg aag ctc   480
Phe Phe Gly Leu Leu Met Val Asn Val Gln Gln Leu Arg Thr Lys Leu
145   150   155   160

aaa ata ttg gac aac act tat gga aat gga act gcc aat aca gca ctc   528
Lys Ile Leu Asp Asn Thr Tyr Gly Asn Gly Thr Ala Asn Thr Ala Leu
            165   170   175

gta tat cac cat gga aaa ctt cta gca tta cag gag gca gat aag ccg   576
Val Tyr His His Gly Lys Leu Leu Ala Leu Gln Glu Ala Asp Lys Pro

                    180   185   190

tac gtc atc aaa gtt ttg gaa gat gga gac ctg caa act ctt ggt ata   624
Tyr Val Ile Lys Val Leu Glu Asp Gly Asp Leu Gln Thr Leu Gly Ile
         195   200   205

ata gat tat gac aag aga ttg acc cac tcc ttc act gct cac cca aaa   672
Ile Asp Tyr Asp Lys Arg Leu Thr His Ser Phe Thr Ala His Pro Lys
         210   215   220

gtt gac ccg gtt acg ggt gaa atg ttt aca ttc ggc tat tcg cat acg   720
Val Asp Pro Val Thr Gly Glu Met Phe Thr Phe Gly Tyr Ser His Thr
225   230   235   240

cca cct tat ctc aca tac aga gtt atc tcg aaa gat ggc att atg cat   768
Pro Pro Tyr Leu Thr Tyr Arg Val Ile Ser Lys Asp Gly Ile Met His
                   245   250   255

gac cca gtc cca att act ata tca gag cct atc atg atg cat gat ttt   816
Asp Pro Val Pro Ile Thr Ile Ser Glu Pro Ile Met Met His Asp Phe
                   260   265   270

gct att act gag act tat gca atc ttc atg gat ctt cct atg cac ttc   864
Ala Ile Thr Glu Thr Tyr Ala Ile Phe Met Asp Leu Pro Met His Phe
         275   280   285

agg cca aag gaa atg gtg aaa gag aag aaa atg ata tac tca ttt gat   912
Arg Pro Lys Glu Met Val Lys Glu Lys Lys Met Ile Tyr Ser Phe Asp
         290   295   300

ccc aca aaa aag gct cgt ttt ggt gtt ctt cca cgc tat gcc aag gat   960
Pro Thr Lys Lys Ala Arg Phe Gly Val Leu Pro Arg Tyr Ala Lys Asp
305   310   315   320

gaa ctt atg att aga tgg ttt gag ctt ccc aac tgc ttt att ttc cac   1008
Glu Leu Met Ile Arg Trp Phe Glu Leu Pro Asn Cys Phe Ile Phe His
                   325   330   335

aac gcc aat gct tgg gaa gaa gag gat gaa gtc gtc ctc atc act tgt   1056
Asn Ala Asn Ala Trp Glu Glu Glu Asp Glu Val Val Leu Ile Thr Cys

                        340   345   350

cgt ctt gag aat cca gat ctt gac atg gtc agt ggg aaa gtg aaa gaa   1104
Arg Leu Glu Asn Pro Asp Leu Asp Met Val Ser Gly Lys Val Lys Glu
        355   360   365

aaa ctc gaa aat ttt ggc aac gaa ctg tac gaa atg aga ttc aac atg   1152
Lys Leu Glu Asn Phe Gly Asn Glu Leu Tyr Glu Met Arg Phe Asn Met
        370   375   380

aaa acg ggc tca gct tct caa aaa aaa cta tcc gca tct gcg gtt gat   1200
Lys Thr Gly Ser Ala Ser Gln Lys Lys Leu Ser Ala Ser Ala Val Asp
385   390   395   400

ttc ccc aga atc aat gag tgc tac acc gga aag aaa cag aga tac gta   1248
Phe Pro Arg Ile Asn Glu Cys Tyr Thr Gly Lys Lys Gln Arg Tyr Val
                405   410   415

tat gga aca att ctg gac agt atc gca aag gtt acc gga atc atc aag   1296
Tyr Gly Thr Ile Leu Asp Ser Ile Ala Lys Val Thr Gly Ile Ile Lys
                420   425   430

ttt gat ctg cat gca gaa gct gag aca ggg aaa aga atg ctg gaa gta   1344
Phe Asp Leu His Ala Glu Ala Glu Thr Gly Lys Arg Met Leu Glu Val
                435   440   445

gga ggt aat atc aaa gga ata tat gac ctg gga gaa ggc aga tat ggt   1392
Gly Gly Asn Ile Lys Gly Ile Tyr Asp Leu Gly Glu Gly Arg Tyr Gly
        450   455   460

tca gag gct atc tat gtt ccg cgt gag aca gca gaa gaa gac gac ggt   1440
Ser Glu Ala Ile Tyr Val Pro Arg Glu Thr Ala Glu Glu Asp Asp Gly
465   470   475   480

tac ttg ata ttc ttt gtt cat gat gaa aac aca ggg aaa tca tgc gtg   1488
Tyr Leu Ile Phe Phe Val His Asp Glu Asn Thr Gly Lys Ser Cys Val
                485   490   495

act gtg ata gac gca aaa aca atg tcg gct gaa ccg gtg gca gtg gtg   1536
Thr Val Ile Asp Ala Lys Thr Met Ser Ala Glu Pro Val Ala Val Val

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           500                    505                    510

gag ctg ccg cac agg gtc cca tat ggc ttc cat gcc ttg ttt gtt aca   1584
Glu Leu Pro His Arg Val Pro Tyr Gly Phe His Ala Leu Phe Val Thr
        515                    520                    525

gag gaa caa ctc cag gaa caa act ctt ata taa                       1617
Glu Glu Gln Leu Gln Glu Gln Thr Leu Ile
        530                    535
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<212> PRT
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Arg Pro Ser Lys Gly Phe Ser Ser Lys Leu Leu Asp Leu Leu Glu Arg
              20                 25                 30

Leu Val Val Lys Leu Met His Asp Ala Ser Leu Pro Leu His Tyr Leu
          35                 40                 45

Ser Gly Asn Phe Ala Pro Ile Arg Asp Glu Thr Pro Pro Val Lys Asp
      50                 55                 60

Leu Pro Val His Gly Phe Leu Pro Glu Cys Leu Asn Gly Glu Phe Val
65                 70                 75                 80

Arg Val Gly Pro Asn Pro Lys Phe Asp Ala Val Ala Gly Tyr His Trp
              85                 90                 95

Phe Asp Gly Asp Gly Met Ile His Gly Val Arg Ile Lys Asp Gly Lys
             100                105                110

Ala Thr Tyr Val Ser Arg Tyr Val Lys Thr Ser Arg Leu Lys Gln Glu
         115                120                125
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Glu Phe Phe Gly Ala Ala Lys Phe Met Lys Ile Gly Asp Leu Lys Gly
    130   135   140

Phe Phe Gly Leu Leu Met Val Asn Val Gln Gln Leu Arg Thr Lys Leu
145   150   155   160

Lys Ile Leu Asp Asn Thr Tyr Gly Asn Gly Thr Ala Asn Thr Ala Leu
                165   170   175

Val Tyr His His Gly Lys Leu Leu Ala Leu Gln Glu Ala Asp Lys Pro
            180   185   190

Tyr Val Ile Lys Val Leu Glu Asp Gly Asp Leu Gln Thr Leu Gly Ile
        195   200   205

Ile Asp Tyr Asp Lys Arg Leu Thr His Ser Phe Thr Ala His Pro Lys
    210   215   220

Val Asp Pro Val Thr Gly Glu Met Phe Thr Phe Gly Tyr Ser His Thr
225   230   235   240

Pro Pro Tyr Leu Thr Tyr Arg Val Ile Ser Lys Asp Gly Ile Met His
            245   250   255

Asp Pro Val Pro Ile Thr Ile Ser Glu Pro Ile Met Met His Asp Phe
            260   265   270

Ala Ile Thr Glu Thr Tyr Ala Ile Phe Met Asp Leu Pro Met His Phe
        275   280   285

Arg Pro Lys Glu Met Val Lys Glu Lys Lys Met Ile Tyr Ser Phe Asp
    290   295   300

Pro Thr Lys Lys Ala Arg Phe Gly Val Leu Pro Arg Tyr Ala Lys Asp
305   310   315   320

Glu Leu Met Ile Arg Trp Phe Glu Leu Pro Asn Cys Phe Ile Phe His
            325   330   335

Asn Ala Asn Ala Trp Glu Glu Glu Asp Glu Val Val Leu Ile Thr Cys
340   345   350

Arg Leu Glu Asn Pro Asp Leu Asp Met Val Ser Gly Lys Val Lys Glu
355   360   365

Lys Leu Glu Asn Phe Gly Asn Glu Leu Tyr Glu Met Arg Phe Asn Met
370   375   380

Lys Thr Gly Ser Ala Ser Gln Lys Lys Leu Ser Ala Ser Ala Val Asp
385   390   395   400

Phe Pro Arg Ile Asn Glu Cys Tyr Thr Gly Lys Lys Gln Arg Tyr Val
405   410   415

Tyr Gly Thr Ile Leu Asp Ser Ile Ala Lys Val Thr Gly Ile Ile Lys
420   425   430

Phe Asp Leu His Ala Glu Ala Glu Thr Gly Lys Arg Met Leu Glu Val
435   440   445

Gly Gly Asn Ile Lys Gly Ile Tyr Asp Leu Gly Glu Gly Arg Tyr Gly
450   455   460

Ser Glu Ala Ile Tyr Val Pro Arg Glu Thr Ala Glu Glu Asp Asp Gly
465   470   475   480

Tyr Leu Ile Phe Phe Val His Asp Glu Asn Thr Gly Lys Ser Cys Val
485   490   495

Thr Val Ile Asp Ala Lys Thr Met Ser Ala Glu Pro Val Ala Val Val
500   505   510

Glu Leu Pro His Arg Val Pro Tyr Gly Phe His Ala Leu Phe Val Thr
515   520   525

Glu Glu Gln Leu Gln Glu Gln Thr Leu Ile
530   535

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Met Gln His Ser Leu Arg Ser Asp Leu Leu Pro Thr Lys Thr Ser Pro
 1               5                   10                  15

cgt tct cat tta ctt cca caa ccc aaa aat gca aat att tct cga cga     96
Arg Ser His Leu Leu Pro Gln Pro Lys Asn Ala Asn Ile Ser Arg Arg
            20                  25                  30

att ctc att aac cct ttc aag ata ccg aca ctt cct gat ctc act tct    144
Ile Leu Ile Asn Pro Phe Lys Ile Pro Thr Leu Pro Asp Leu Thr Ser
            35                  40                  45

ccg gtt ccg tca ccg gtt aag ctc aaa cca acg tat cca aac tta aac    192
Pro Val Pro Ser Pro Val Lys Leu Lys Pro Thr Tyr Pro Asn Leu Asn
        50                  55                  60

ctt ctt cag aag cta gcg gct acg atg ctc gac aag att gag tcc tct    240
Leu Leu Gln Lys Leu Ala Ala Thr Met Leu Asp Lys Ile Glu Ser Ser
65                  70                  75                  80

atc gtt att cct atg gag cag aat cgc ccg ctt cct aaa ccg acc gac    288
Ile Val Ile Pro Met Glu Gln Asn Arg Pro Leu Pro Lys Pro Thr Asp
                85                  90                  95

ccg gcg gtt caa tta tca ggt aac ttc gct ccg gtt aat gaa tgt ccg    336
Pro Ala Val Gln Leu Ser Gly Asn Phe Ala Pro Val Asn Glu Cys Pro
            100                 105                 110

gtt cag aac ggt tta gaa gtg gtt ggt cag att cct tct tgt cta aaa    384
Val Gln Asn Gly Leu Glu Val Val Gly Gln Ile Pro Ser Cys Leu Lys
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115   120   125

gga gtt tac atc cgt aac ggt gca aac cct atg ttt ccg ccg tta gcc   432
Gly Val Tyr Ile Arg Asn Gly Ala Asn Pro Met Phe Pro Pro Leu Ala
   130   135   140

gga cat cat tta ttt gac ggt gac gga atg att cac gcc gtt agt atc   480
Gly His His Leu Phe Asp Gly Asp Gly Met Ile His Ala Val Ser Ile
145   150   155   160

ggt ttt gat aac cag gtt agt tac agc tgc cgg tac act aaa aca aac   528
Gly Phe Asp Asn Gln Val Ser Tyr Ser Cys Arg Tyr Thr Lys Thr Asn
                165   170   175

cgg ctt gtt caa gaa acc gcg ctt gga cga tcg gtt ttc cct aaa cca   576
Arg Leu Val Gln Glu Thr Ala Leu Gly Arg Ser Val Phe Pro Lys Pro
            180   185   190

atc ggc gag ctt cac ggc cat tcc ggt cta gct cga ctc gct ctc ttc   624
Ile Gly Glu Leu His Gly His Ser Gly Leu Ala Arg Leu Ala Leu Phe
         195   200   205

acg gct cga gct ggg atc ggt cta gtg gac ggg aca cgt ggc atg ggc   672
Thr Ala Arg Ala Gly Ile Gly Leu Val Asp Gly Thr Arg Gly Met Gly
   210   215   220

gta gct aac gcc ggt gtg gtt ttc ttt aac ggc agg tta tta gcc atg   720
Val Ala Asn Ala Gly Val Val Phe Phe Asn Gly Arg Leu Leu Ala Met
225   230   235   240

tca gaa gat gat ctt cct tac caa gtg aag atc gac ggt caa gga gat   768
Ser Glu Asp Asp Leu Pro Tyr Gln Val Lys Ile Asp Gly Gln Gly Asp
                245   250   255

ctt gag acg atc gga cgg ttc gga ttc gat gac cag att gac tct tca   816
Leu Glu Thr Ile Gly Arg Phe Gly Phe Asp Asp Gln Ile Asp Ser Ser
            260   265   270

gtg ata gcg cat cct aag gtg gac gcg acc aca gga gat ctc cat aca   864
Val Ile Ala His Pro Lys Val Asp Ala Thr Thr Gly Asp Leu His Thr

275   280   285

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ctg agc tac aac gtt ttg aag aaa cct cat ctc agg tat ctt aaa ttc    912
Leu Ser Tyr Asn Val Leu Lys Lys Pro His Leu Arg Tyr Leu Lys Phe
    290             295             300

aac acg tgc ggg aaa aag aca cgt gac gtg gag atc acg ctc cct gaa    960
Asn Thr Cys Gly Lys Lys Thr Arg Asp Val Glu Ile Thr Leu Pro Glu
305             310             315             320

cca acg atg att cat gat ttc gcg ata acc gag aat ttt gtc gtt ata    1008
Pro Thr Met Ile His Asp Phe Ala Ile Thr Glu Asn Phe Val Val Ile
                325             330             335

ccg gat cag caa atg gta ttc aaa tta tcc gaa atg att cgg ggc ggg    1056
Pro Asp Gln Gln Met Val Phe Lys Leu Ser Glu Met Ile Arg Gly Gly
            340             345             350

tca ccc gtt atc tac gtt aaa gaa aaa atg gcg aga ttt gga gtt ttg    1104
Ser Pro Val Ile Tyr Val Lys Glu Lys Met Ala Arg Phe Gly Val Leu
            355             360             365

tca aag cag gat ctg acc ggg tcg gat ata aat tgg gtt gat gta ccg    1152
Ser Lys Gln Asp Leu Thr Gly Ser Asp Ile Asn Trp Val Asp Val Pro
    370             375             380

gat tgt ttc tgt ttc cat cta tgg aat gcg tgg gaa gag aga acc gaa    1200
Asp Cys Phe Cys Phe His Leu Trp Asn Ala Trp Glu Glu Arg Thr Glu
385             390             395             400

gag gga gac cca gtt atc gtc gta atc ggg tca tgt atg agc cca ccc    1248
Glu Gly Asp Pro Val Ile Val Val Ile Gly Ser Cys Met Ser Pro Pro
                405             410             415

gac acg atc ttt agt gaa tca gga gaa cca acc cgg gtt gaa tta agt    1296
Asp Thr Ile Phe Ser Glu Ser Gly Glu Pro Thr Arg Val Glu Leu Ser
            420             425             430

gag atc cgg tta aac atg cgt aca aaa gaa tcg aac cgt aag gtt atc    1344
Glu Ile Arg Leu Asn Met Arg Thr Lys Glu Ser Asn Arg Lys Val Ile
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                435                 440                 445


gta acc gga gtg aat tta gaa gcg ggt cac ata aac cgt agt tac gtg    1392
Val Thr Gly Val Asn Leu Glu Ala Gly His Ile Asn Arg Ser Tyr Val
    450                 455                 460


ggc cgg aaa agc cag ttc gtt tac ata gca ata gcc gat cct tgg ccc    1440
Gly Arg Lys Ser Gln Phe Val Tyr Ile Ala Ile Ala Asp Pro Trp Pro
465                 470                 475                 480


aaa tgc agt ggc att gcg aag gta gat ata caa aac ggc acc gtt tca    1488
Lys Cys Ser Gly Ile Ala Lys Val Asp Ile Gln Asn Gly Thr Val Ser
                485                 490                 495


gag ttt aat tac gga ccg agc cgg ttc ggt gga gaa ccg tgc ttt gta    1536
Glu Phe Asn Tyr Gly Pro Ser Arg Phe Gly Gly Glu Pro Cys Phe Val
                500                 505                 510


ccg gag gga gaa gga gaa gaa gac aaa ggt tat gta atg ggg ttt gtg    1584
Pro Glu Gly Glu Gly Glu Glu Asp Lys Gly Tyr Val Met Gly Phe Val
                515                 520                 525


aga gac gaa gag aaa gac gag tcg gag ttt gtg gtg gtc gac gcg acg    1632
Arg Asp Glu Glu Lys Asp Glu Ser Glu Phe Val Val Val Asp Ala Thr
    530                 535                 540


gat atg aag caa gtc gcg gcg gtg cgc ttg ccg gag agg gta cct tat    1680
Asp Met Lys Gln Val Ala Ala Val Arg Leu Pro Glu Arg Val Pro Tyr
545                 550                 555                 560


ggt ttc cat gga acg ttc gtg agc gag aat cag ttg aag gaa caa gtt    1728
Gly Phe His Gly Thr Phe Val Ser Glu Asn Gln Leu Lys Glu Gln Val
                565                 570                 575


ttc tga                                                            1734
Phe
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<210> 10
<211> 577

<212> PRT
<213> Arabidopsis thaliana

<400> 10
Met Gln His Ser Leu Arg Ser Asp Leu Leu Pro Thr Lys Thr Ser Pro
1   5   10   15

Arg Ser His Leu Leu Pro Gln Pro Lys Asn Ala Asn Ile Ser Arg Arg
            20   25   30

Ile Leu Ile Asn Pro Phe Lys Ile Pro Thr Leu Pro Asp Leu Thr Ser
        35   40   45

Pro Val Pro Ser Pro Val Lys Leu Lys Pro Thr Tyr Pro Asn Leu Asn
        50   55   60

Leu Leu Gln Lys Leu Ala Ala Thr Met Leu Asp Lys Ile Glu Ser Ser
65   70   75   80

Ile Val Ile Pro Met Glu Gln Asn Arg Pro Leu Pro Lys Pro Thr Asp
            85   90   95

Pro Ala Val Gln Leu Ser Gly Asn Phe Ala Pro Val Asn Glu Cys Pro
        100   105   110

Val Gln Asn Gly Leu Glu Val Val Gly Gln Ile Pro Ser Cys Leu Lys
        115   120   125

Gly Val Tyr Ile Arg Asn Gly Ala Asn Pro Met Phe Pro Pro Leu Ala
    130   135   140

Gly His His Leu Phe Asp Gly Asp Gly Met Ile His Ala Val Ser Ile
145   150   155   160

Gly Phe Asp Asn Gln Val Ser Tyr Ser Cys Arg Tyr Thr Lys Thr Asn
            165   170   175

Arg Leu Val Gln Glu Thr Ala Leu Gly Arg Ser Val Phe Pro Lys Pro
        180   185   190

Ile Gly Glu Leu His Gly His Ser Gly Leu Ala Arg Leu Ala Leu Phe
        195   200   205

Thr Ala Arg Ala Gly Ile Gly Leu Val Asp Gly Thr Arg Gly Met Gly
        210   215   220

Val Ala Asn Ala Gly Val Val Phe Phe Asn Gly Arg Leu Leu Ala Met
225   230   235   240

Ser Glu Asp Asp Leu Pro Tyr Gln Val Lys Ile Asp Gly Gln Gly Asp
                245   250   255

Leu Glu Thr Ile Gly Arg Phe Gly Phe Asp Asp Gln Ile Asp Ser Ser
                260   265   270

Val Ile Ala His Pro Lys Val Asp Ala Thr Thr Gly Asp Leu His Thr
        275   280   285

Leu Ser Tyr Asn Val Leu Lys Lys Pro His Leu Arg Tyr Leu Lys Phe
        290   295   300

Asn Thr Cys Gly Lys Lys Thr Arg Asp Val Glu Ile Thr Leu Pro Glu
305   310   315   320

Pro Thr Met Ile His Asp Phe Ala Ile Thr Glu Asn Phe Val Val Ile
                325   330   335

Pro Asp Gln Gln Met Val Phe Lys Leu Ser Glu Met Ile Arg Gly Gly
                340   345   350

Ser Pro Val Ile Tyr Val Lys Glu Lys Met Ala Arg Phe Gly Val Leu
        355   360   365

Ser Lys Gln Asp Leu Thr Gly Ser Asp Ile Asn Trp Val Asp Val Pro
        370   375   380

Asp Cys Phe Cys Phe His Leu Trp Asn Ala Trp Glu Glu Arg Thr Glu
385   390   395   400

Glu Gly Asp Pro Val Ile Val Val Ile Gly Ser Cys Met Ser Pro Pro

                405   410   415

Asp Thr Ile Phe Ser Glu Ser Gly Glu Pro Thr Arg Val Glu Leu Ser
         420   425   430

Glu Ile Arg Leu Asn Met Arg Thr Lys Glu Ser Asn Arg Lys Val Ile
         435   440   445

Val Thr Gly Val Asn Leu Glu Ala Gly His Ile Asn Arg Ser Tyr Val
         450   455   460

Gly Arg Lys Ser Gln Phe Val Tyr Ile Ala Ile Ala Asp Pro Trp Pro
465   470   475   480

Lys Cys Ser Gly Ile Ala Lys Val Asp Ile Gln Asn Gly Thr Val Ser
             485   490   495

Glu Phe Asn Tyr Gly Pro Ser Arg Phe Gly Gly Glu Pro Cys Phe Val
             500   505   510

Pro Glu Gly Glu Gly Glu Glu Asp Lys Gly Tyr Val Met Gly Phe Val
         515   520   525

Arg Asp Glu Glu Lys Asp Glu Ser Glu Phe Val Val Val Asp Ala Thr
         530   535   540

Asp Met Lys Gln Val Ala Ala Val Arg Leu Pro Glu Arg Val Pro Tyr
545   550   555   560

Gly Phe His Gly Thr Phe Val Ser Glu Asn Gln Leu Lys Glu Gln Val
             565   570   575

Phe

<210> 11
<211> 1839
<212> DNA
<213> Vigna unguiculata

<220>
<221> CDS
<222> (1)..(1839)

<400> 11
atg cct tca tca gct tca aac act tgg ttt aac gcc aca ctc cca tct   48
Met Pro Ser Ser Ala Ser Asn Thr Trp Phe Asn Ala Thr Leu Pro Ser
1   5   10   15

ccc ccc ttc aaa gac cta cct tcc aca tct tct ccc aca aac tta ctt   96
Pro Pro Phe Lys Asp Leu Pro Ser Thr Ser Ser Pro Thr Asn Leu Leu
            20   25   30

cct tta agg aaa aca tcc tct tcc aac acc atc aca tgt tcc ctt caa   144
Pro Leu Arg Lys Thr Ser Ser Ser Asn Thr Ile Thr Cys Ser Leu Gln
            35   40   45

aca ctc cac ttc ccc aaa cag tac caa cca aca tcc aca tcc aca tcc   192
Thr Leu His Phe Pro Lys Gln Tyr Gln Pro Thr Ser Thr Ser Thr Ser
    50   55   60

aca gcc acc acc aca aca ccc acc cca atc aaa act acc acc atc acc   240
Thr Ala Thr Thr Thr Thr Pro Thr Pro Ile Lys Thr Thr Thr Ile Thr
65   70   75   80

acc acc aca ccg ccc agg gaa acc aac cct ctc tct gac acc aac caa   288
Thr Thr Thr Pro Pro Arg Glu Thr Asn Pro Leu Ser Asp Thr Asn Gln
                85   90   95

cca tta cct caa aaa tgg aac ttt ctc cag aaa gcc gct gcc acg gcc   336
Pro Leu Pro Gln Lys Trp Asn Phe Leu Gln Lys Ala Ala Ala Thr Ala
            100   105   110

ttg gac ctg gtc gaa acg gcg ctc gtc tcg cac gag cgc aaa cac ccg   384
Leu Asp Leu Val Glu Thr Ala Leu Val Ser His Glu Arg Lys His Pro
            115   120   125

ctc ccc aaa acg gcg gac ccg agg gtc caa atc gcc ggg aac ttc gcg   432
Leu Pro Lys Thr Ala Asp Pro Arg Val Gln Ile Ala Gly Asn Phe Ala
        130   135   140

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ccg gtg ccg gag cat gcc gcg gat caa gga ctc ccg gtg gtc gga aaa    480
Pro Val Pro Glu His Ala Ala Asp Gln Gly Leu Pro Val Val Gly Lys
145             150             155             160

atc ccc aaa tgc att gac ggc gtg tac gtg cgc aac ggt gcc aat ccg    528
Ile Pro Lys Cys Ile Asp Gly Val Tyr Val Arg Asn Gly Ala Asn Pro
            165             170             175

ctc tac gag cct gtg gcc ggg cac cac ttc ttc gac ggc gac ggc atg    576
Leu Tyr Glu Pro Val Ala Gly His His Phe Phe Asp Gly Asp Gly Met
            180             185             190

gtc cac gcc gtg aag ttc acg aac ggc gcc gcc agc tac gcc tgc cgc    624
Val His Ala Val Lys Phe Thr Asn Gly Ala Ala Ser Tyr Ala Cys Arg
            195             200             205

ttc acc gag acg cag cgt ctc tcg cag gag aaa tct cta ggc cgc ccg    672
Phe Thr Glu Thr Gln Arg Leu Ser Gln Glu Lys Ser Leu Gly Arg Pro
    210             215             220

gtg ttc ccg aag gcc atc ggg gag ctc cac ggc cac tcc ggc atc gcg    720
Val Phe Pro Lys Ala Ile Gly Glu Leu His Gly His Ser Gly Ile Ala
225             230             235             240

cgg ctc ctc ctc ttc tac gcg cgc ggt ctc ttc ggg ctc gtt gat ggg    768
Arg Leu Leu Leu Phe Tyr Ala Arg Gly Leu Phe Gly Leu Val Asp Gly
                245             250             255

tcc cag ggc atg ggc gtg gcg aac gcc ggt ctc gtc tac ttc aac aac    816
Ser Gln Gly Met Gly Val Ala Asn Ala Gly Leu Val Tyr Phe Asn Asn
            260             265             270

cac ctc ttg gcc atg tcc gaa gac gat tta ccc tac cac gtg aga atc    864
His Leu Leu Ala Met Ser Glu Asp Asp Leu Pro Tyr His Val Arg Ile
            275             280             285

acc cct aac ggc gac tta acc acc gtt ggc cgt tac gac ttc aac ggg    912
Thr Pro Asn Gly Asp Leu Thr Thr Val Gly Arg Tyr Asp Phe Asn Gly
        290             295             300
```

cag ctc aac tca aca atg atc gcc cac ccg aaa ctg gac ccc gtc gac   960
Gln Leu Asn Ser Thr Met Ile Ala His Pro Lys Leu Asp Pro Val Asp
305   310   315   320

ggc gac ctc cac gcg ctc agc tac gac gtc att cag aag cct tac ctc   1008
Gly Asp Leu His Ala Leu Ser Tyr Asp Val Ile Gln Lys Pro Tyr Leu
           325   330   335

aag tac ttc cgt ttc tcc ccc gac ggc gtc aag tcc ccc gac gtg gaa   1056
Lys Tyr Phe Arg Phe Ser Pro Asp Gly Val Lys Ser Pro Asp Val Glu
           340   345   350

atc ccc ctg aag gag ccc acc atg atg cac gat ttc gcc ata acg gag   1104
Ile Pro Leu Lys Glu Pro Thr Met Met His Asp Phe Ala Ile Thr Glu
           355   360   365

aat ttc gtc gtc gtc ccc gac cag cag gtg gtc ttc aaa cta acg gag   1152
Asn Phe Val Val Val Pro Asp Gln Gln Val Val Phe Lys Leu Thr Glu
370   375   380

atg atc acc ggc ggg tcc ccc gtg gtc tac gac aag aac aaa acc tca   1200
Met Ile Thr Gly Gly Ser Pro Val Val Tyr Asp Lys Asn Lys Thr Ser
385   390   395   400

cgg ttt ggg att ctg cac aag aat gcg aag gac gcg aat gcg atg cgg   1248
Arg Phe Gly Ile Leu His Lys Asn Ala Lys Asp Ala Asn Ala Met Arg
           405   410   415

tgg atc gac gcg ccg gat tgt ttc tgc ttc cac ctc tgg aac gcg tgg   1296
Trp Ile Asp Ala Pro Asp Cys Phe Cys Phe His Leu Trp Asn Ala Trp
           420   425   430

gag gag ccc gaa acc gag gag gtt gtg gtg att ggg tcc tgc atg acc   1344
Glu Glu Pro Glu Thr Glu Glu Val Val Val Ile Gly Ser Cys Met Thr
           435   440   445

cct gcg gac tcc att ttc aac gaa tgc gag gag agt ttg aag agc gtg   1392
Pro Ala Asp Ser Ile Phe Asn Glu Cys Glu Glu Ser Leu Lys Ser Val
           450   455   460

```
ctg tca gag ata agg ctg aac ttg agg acc ggc aag tcc act cgg cgc    1440
Leu Ser Glu Ile Arg Leu Asn Leu Arg Thr Gly Lys Ser Thr Arg Arg
465             470             475             480


ccc att atc tcc gac gcc gaa caa gtg aac ctg gaa gcc ggc atg gtg    1488
Pro Ile Ile Ser Asp Ala Glu Gln Val Asn Leu Glu Ala Gly Met Val
            485             490             495


aac aga aac aag ctc gga agg aag acc cag ttc gcg tat ctg gct ctg    1536
Asn Arg Asn Lys Leu Gly Arg Lys Thr Gln Phe Ala Tyr Leu Ala Leu
            500             505             510


gcg gag ccc tgg ccc aaa gtc tcg ggc ttt gcg aaa gtt gat ttg ctg    1584
Ala Glu Pro Trp Pro Lys Val Ser Gly Phe Ala Lys Val Asp Leu Leu
        515             520             525


agt ggg gaa gtg aag aag tac atg tat gga gaa gag aag ttc ggt ggg    1632
Ser Gly Glu Val Lys Lys Tyr Met Tyr Gly Glu Glu Lys Phe Gly Gly
        530             535             540


gag cct ctg ttt ctt ccc aac ggc caa aaa gaa gac gat ggg tat att    1680
Glu Pro Leu Phe Leu Pro Asn Gly Gln Lys Glu Asp Asp Gly Tyr Ile
545             550             555             560


ctg gca ttc gtg cac gac gag aaa gaa tgg aaa tcc gag ctg cag att    1728
Leu Ala Phe Val His Asp Glu Lys Glu Trp Lys Ser Glu Leu Gln Ile
            565             570             575


gtg aat gcc caa aat tta aag ctc gaa gct tcc atc aaa ctc ccc tct    1776
Val Asn Ala Gln Asn Leu Lys Leu Glu Ala Ser Ile Lys Leu Pro Ser
            580             585             590


cgt gtt ccc tac ggt ttt cat gga act ttc att cat tcc aag gat ttg    1824
Arg Val Pro Tyr Gly Phe His Gly Thr Phe Ile His Ser Lys Asp Leu
            595             600             605


agg aaa caa gct tga                                                1839
Arg Lys Gln Ala
            610
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<210> 12
<211> 612
<212> PRT
<213> Vigna unguiculata

<400> 12
Met Pro Ser Ser Ala Ser Asn Thr Trp Phe Asn Ala Thr Leu Pro Ser
1   5   10   15

Pro Pro Phe Lys Asp Leu Pro Ser Thr Ser Ser Pro Thr Asn Leu Leu
            20   25   30

Pro Leu Arg Lys Thr Ser Ser Ser Asn Thr Ile Thr Cys Ser Leu Gln
            35   40   45

Thr Leu His Phe Pro Lys Gln Tyr Gln Pro Thr Ser Thr Ser Thr Ser
        50   55   60

Thr Ala Thr Thr Thr Thr Pro Thr Pro Ile Lys Thr Thr Thr Ile Thr
65   70   75   80

Thr Thr Thr Pro Pro Arg Glu Thr Asn Pro Leu Ser Asp Thr Asn Gln
                85   90   95

Pro Leu Pro Gln Lys Trp Asn Phe Leu Gln Lys Ala Ala Ala Thr Ala
            100   105   110

Leu Asp Leu Val Glu Thr Ala Leu Val Ser His Glu Arg Lys His Pro
            115   120   125

Leu Pro Lys Thr Ala Asp Pro Arg Val Gln Ile Ala Gly Asn Phe Ala
        130   135   140

Pro Val Pro Glu His Ala Ala Asp Gln Gly Leu Pro Val Val Gly Lys
145   150   155   160

Ile Pro Lys Cys Ile Asp Gly Val Tyr Val Arg Asn Gly Ala Asn Pro
                165   170   175

Leu Tyr Glu Pro Val Ala Gly His His Phe Phe Asp Gly Asp Gly Met
         180   185   190

Val His Ala Val Lys Phe Thr Asn Gly Ala Ala Ser Tyr Ala Cys Arg
         195   200   205

Phe Thr Glu Thr Gln Arg Leu Ser Gln Glu Lys Ser Leu Gly Arg Pro
         210   215   220

Val Phe Pro Lys Ala Ile Gly Glu Leu His Gly His Ser Gly Ile Ala
225   230   235   240

Arg Leu Leu Leu Phe Tyr Ala Arg Gly Leu Phe Gly Leu Val Asp Gly
               245   250   255

Ser Gln Gly Met Gly Val Ala Asn Ala Gly Leu Val Tyr Phe Asn Asn
               260   265   270

His Leu Leu Ala Met Ser Glu Asp Asp Leu Pro Tyr His Val Arg Ile
         275   280   285

Thr Pro Asn Gly Asp Leu Thr Thr Val Gly Arg Tyr Asp Phe Asn Gly
         290   295   300

Gln Leu Asn Ser Thr Met Ile Ala His Pro Lys Leu Asp Pro Val Asp
305   310   315   320

Gly Asp Leu His Ala Leu Ser Tyr Asp Val Ile Gln Lys Pro Tyr Leu
               325   330   335

Lys Tyr Phe Arg Phe Ser Pro Asp Gly Val Lys Ser Pro Asp Val Glu
               340   345   350

Ile Pro Leu Lys Glu Pro Thr Met Met His Asp Phe Ala Ile Thr Glu
               355   360   365

Asn Phe Val Val Val Pro Asp Gln Gln Val Val Phe Lys Leu Thr Glu
         370   375   380

Met Ile Thr Gly Gly Ser Pro Val Val Tyr Asp Lys Asn Lys Thr Ser
385   390   395   400

Arg Phe Gly Ile Leu His Lys Asn Ala Lys Asp Ala Asn Ala Met Arg
              405   410   415

Trp Ile Asp Ala Pro Asp Cys Phe Cys Phe His Leu Trp Asn Ala Trp
          420   425   430

Glu Glu Pro Glu Thr Glu Glu Val Val Val Ile Gly Ser Cys Met Thr
          435   440   445

Pro Ala Asp Ser Ile Phe Asn Glu Cys Glu Glu Ser Leu Lys Ser Val
          450   455   460

Leu Ser Glu Ile Arg Leu Asn Leu Arg Thr Gly Lys Ser Thr Arg Arg
465   470   475   480

Pro Ile Ile Ser Asp Ala Glu Gln Val Asn Leu Glu Ala Gly Met Val
              485   490   495

Asn Arg Asn Lys Leu Gly Arg Lys Thr Gln Phe Ala Tyr Leu Ala Leu
          500   505   510

Ala Glu Pro Trp Pro Lys Val Ser Gly Phe Ala Lys Val Asp Leu Leu
          515   520   525

Ser Gly Glu Val Lys Lys Tyr Met Tyr Gly Glu Glu Lys Phe Gly Gly
          530   535   540

Glu Pro Leu Phe Leu Pro Asn Gly Gln Lys Glu Asp Asp Gly Tyr Ile
545   550   555   560

Leu Ala Phe Val His Asp Glu Lys Glu Trp Lys Ser Glu Leu Gln Ile
              565   570   575

Val Asn Ala Gln Asn Leu Lys Leu Glu Ala Ser Ile Lys Leu Pro Ser
              580   585   590

Arg Val Pro Tyr Gly Phe His Gly Thr Phe Ile His Ser Lys Asp Leu

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              595              600              605

Arg Lys Gln Ala
        610


<210> 13
<211> 1815
<212> DNA
<213> Zea mays

<220>
<221> CDS
<222> (1)..(1815)

<400> 13
atg cag ggt ctc gcc ccg ccc acc tct gtt tcc ata cac cgg cac ctg    48
Met Gln Gly Leu Ala Pro Pro Thr Ser Val Ser Ile His Arg His Leu
  1               5                  10                  15


ccg gcc cgg tcc agg gcc cgg gcc tcc aat tcc gtc agg ttc tcg ccg    96
Pro Ala Arg Ser Arg Ala Arg Ala Ser Asn Ser Val Arg Phe Ser Pro
            20                  25                  30


cgc gcc gtc agc tcc gtg ccg ccc gcc gag tgc ctc cag gcg ccg ttc   144
Arg Ala Val Ser Ser Val Pro Pro Ala Glu Cys Leu Gln Ala Pro Phe
            35                  40                  45


cac aag ccc gtc gcc gac ctg cct gcg ccg tcc agg aag ccc gcc gcc   192
His Lys Pro Val Ala Asp Leu Pro Ala Pro Ser Arg Lys Pro Ala Ala
        50                  55                  60


att gcc gtc cca ggg cac gcc gcg gcg ccg agg aaa gcg gag ggc ggc   240
Ile Ala Val Pro Gly His Ala Ala Ala Pro Arg Lys Ala Glu Gly Gly
 65                  70                  75                  80


aag aag cag ctc aac ttg ttc cag cgc gcc gcg gcg gcc gcg ctc gac   288
Lys Lys Gln Leu Asn Leu Phe Gln Arg Ala Ala Ala Ala Ala Leu Asp
                85                  90                  95
```

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gcg ttc gag gaa ggg ttc gtg gcc aac gtc ctc gag cgg ccc cac ggg      336
Ala Phe Glu Glu Gly Phe Val Ala Asn Val Leu Glu Arg Pro His Gly
            100             105             110

ctg ccc agc acg gcc gac ccg gcc gtg cag atc gcc ggc aac ttc gcg      384
Leu Pro Ser Thr Ala Asp Pro Ala Val Gln Ile Ala Gly Asn Phe Ala
            115             120             125

ccc gtc ggg gag agg ccg ccc gtg cac gag ctc ccc gtc tcc ggc cgc      432
Pro Val Gly Glu Arg Pro Pro Val His Glu Leu Pro Val Ser Gly Arg
            130             135             140

atc ccg ccc ttc atc gac ggg gtc tac gcg cgc aac ggc gcc aac ccc      480
Ile Pro Pro Phe Ile Asp Gly Val Tyr Ala Arg Asn Gly Ala Asn Pro
145             150             155             160

tgc ttc gac ccc gtc gcg ggg cac cac ctc ttc gac ggc gac ggc atg      528
Cys Phe Asp Pro Val Ala Gly His His Leu Phe Asp Gly Asp Gly Met
                165             170             175

gtg cac gcg ctg cgg ata cgc aac ggc gcc gcc gag tcc tac gcc tgc      576
Val His Ala Leu Arg Ile Arg Asn Gly Ala Ala Glu Ser Tyr Ala Cys
                180             185             190

cgc ttc acg gag acc gcg cgc ctg cgc cag gag cgc gcg atc ggc cgc      624
Arg Phe Thr Glu Thr Ala Arg Leu Arg Gln Glu Arg Ala Ile Gly Arg
            195             200             205

ccc gtc ttc ccc aag gcc att ggc gag ctg cac ggg cac tcc ggg atc      672
Pro Val Phe Pro Lys Ala Ile Gly Glu Leu His Gly His Ser Gly Ile
            210             215             220

gcg cgc ctc gcc ctg ttc tac gcg cgc gcc gcg tgc ggc ctc gtg gac      720
Ala Arg Leu Ala Leu Phe Tyr Ala Arg Ala Ala Cys Gly Leu Val Asp
225             230             235             240

ccc tcg gcc ggc acc ggc gtg gcc aac gcc ggc ctc gtc tac ttc aac      768
Pro Ser Ala Gly Thr Gly Val Ala Asn Ala Gly Leu Val Tyr Phe Asn
                245             250             255
```

ggc cgc ctg ctc gcc atg tcc gag gac gac ctc ccc tac cac gtc cgc   816
Gly Arg Leu Leu Ala Met Ser Glu Asp Asp Leu Pro Tyr His Val Arg
          260   265   270

gtg gcg gac gac ggc gac ctc gag acc gtc ggc cgc tac gac ttc gac   864
Val Ala Asp Asp Gly Asp Leu Glu Thr Val Gly Arg Tyr Asp Phe Asp
          275   280   285

ggg cag ctc ggc tgc gcc atg atc gcg cac ccc aag ctg gac ccg gcc   912
Gly Gln Leu Gly Cys Ala Met Ile Ala His Pro Lys Leu Asp Pro Ala
          290   295   300

acc ggg gag ctc cac gcg ctc agc tac gac gtc atc aag agg ccg tac   960
Thr Gly Glu Leu His Ala Leu Ser Tyr Asp Val Ile Lys Arg Pro Tyr
305   310   315   320

ctc aag tac ttc tac ttc agg ccc gac ggc acc aag tcc gac gac gtg   1008
Leu Lys Tyr Phe Tyr Phe Arg Pro Asp Gly Thr Lys Ser Asp Asp Val
          325   330   335

gag atc ccg ctg gag cag ccc acg atg atc cac gac ttc gcc atc acc   1056
Glu Ile Pro Leu Glu Gln Pro Thr Met Ile His Asp Phe Ala Ile Thr
          340   345   350

gag aac ttc gtg gtt gtg ccc gac cac cag gtg gtg ttc aag ctc cag   1104
Glu Asn Phe Val Val Val Pro Asp His Gln Val Val Phe Lys Leu Gln
          355   360   365

gag atg ctg cgc ggc ggg tcg ccc gtg gtg ctg gac aag gag aag acg   1152
Glu Met Leu Arg Gly Gly Ser Pro Val Val Leu Asp Lys Glu Lys Thr
          370   375   380

tcg cgg ttc ggc gtg ctc ccc aag cac gcc gcg gac gcg tcg gag atg   1200
Ser Arg Phe Gly Val Leu Pro Lys His Ala Ala Asp Ala Ser Glu Met
385   390   395   400

gcg tgg gtg gac gtg ccg gac tgc ttc tgc ttc cac ctg tgg aac gcg   1248
Ala Trp Val Asp Val Pro Asp Cys Phe Cys Phe His Leu Trp Asn Ala
          405   410   415

tgg gag gac gag gcg acg ggc gag gtg gtg gtg atc ggc tcc tgc atg   1296
Trp Glu Asp Glu Ala Thr Gly Glu Val Val Val Ile Gly Ser Cys Met
         420   425   430

acc ccc gcc gac tcc atc ttc aac gag tcc gac gag cgc ctg gag agc   1344
Thr Pro Ala Asp Ser Ile Phe Asn Glu Ser Asp Glu Arg Leu Glu Ser
         435   440   445

gtg ctg acc gag atc cgc ctg gac gcg cgc acg ggc cgg tcc acg cgc   1392
Val Leu Thr Glu Ile Arg Leu Asp Ala Arg Thr Gly Arg Ser Thr Arg
         450   455   460

cgc gcc gtc ctg ccg ccg tcg cag cag gag aac ctg gag gtg ggc atg   1440
Arg Ala Val Leu Pro Pro Ser Gln Gln Glu Asn Leu Glu Val Gly Met
465   470   475   480

gtg aac cgc aac ctg ctg ggc cgc gag agc cgg tac gcg tac ctc gcg   1488
Val Asn Arg Asn Leu Leu Gly Arg Glu Ser Arg Tyr Ala Tyr Leu Ala
              485   490   495

gtg gcg gag ccg tgg ccc aag gag tcg ggc ttc gcc aag gag gac ctg   1536
Val Ala Glu Pro Trp Pro Lys Glu Ser Gly Phe Ala Lys Glu Asp Leu
         500   505   510

tcc acg ggc gag ctc acc aag ttc gag tac ggc gag ggc cgg ttc ggc   1584
Ser Thr Gly Glu Leu Thr Lys Phe Glu Tyr Gly Glu Gly Arg Phe Gly
         515   520   525

ggc gag ccc tgc ttc gtt ccc atg gac ccg gcc gcg gcc cac ccg cgc   1632
Gly Glu Pro Cys Phe Val Pro Met Asp Pro Ala Ala Ala His Pro Arg
         530   535   540

ggc gag gac gac ggg tac gtg ctc acc ttc gtc cac gac gag cgc gcc   1680
Gly Glu Asp Asp Gly Tyr Val Leu Thr Phe Val His Asp Glu Arg Ala
545   550   555   560

ggc acg tcg gag cta ctt gtg gtc aat gcc gcc gac atc cgg ctg gag   1728
Gly Thr Ser Glu Leu Leu Val Val Asn Ala Ala Asp Ile Arg Leu Glu
         565   570   575

```
gcc acg gtt cag ctg ccg tcc cgc gtg ccc ttc ggc ttc cac ggc acc      1776
Ala Thr Val Gln Leu Pro Ser Arg Val Pro Phe Gly Phe His Gly Thr
          580                 585                 590

ttc atc acg ggc cag gag ctc gag gcc cag gcg gcc tga                   1815
Phe Ile Thr Gly Gln Glu Leu Glu Ala Gln Ala Ala
          595                 600                 605
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<210> 14
<211> 604
<212> PRT
<213> Zea mays

<400> 14

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Met Gln Gly Leu Ala Pro Pro Thr Ser Val Ser Ile His Arg His Leu
  1               5                  10                  15

Pro Ala Arg Ser Arg Ala Arg Ala Ser Asn Ser Val Arg Phe Ser Pro
              20                  25                  30

Arg Ala Val Ser Ser Val Pro Pro Ala Glu Cys Leu Gln Ala Pro Phe
          35                  40                  45

His Lys Pro Val Ala Asp Leu Pro Ala Pro Ser Arg Lys Pro Ala Ala
      50                  55                  60

Ile Ala Val Pro Gly His Ala Ala Ala Pro Arg Lys Ala Glu Gly Gly
 65                  70                  75                  80

Lys Lys Gln Leu Asn Leu Phe Gln Arg Ala Ala Ala Ala Ala Leu Asp
              85                  90                  95

Ala Phe Glu Glu Gly Phe Val Ala Asn Val Leu Glu Arg Pro His Gly
              100                 105                 110

Leu Pro Ser Thr Ala Asp Pro Ala Val Gln Ile Ala Gly Asn Phe Ala
          115                 120                 125

Pro Val Gly Glu Arg Pro Pro Val His Glu Leu Pro Val Ser Gly Arg
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130   135   140

Ile Pro Pro Phe Ile Asp Gly Val Tyr Ala Arg Asn Gly Ala Asn Pro
145   150   155   160

Cys Phe Asp Pro Val Ala Gly His His Leu Phe Asp Gly Asp Gly Met
                 165   170   175

Val His Ala Leu Arg Ile Arg Asn Gly Ala Ala Glu Ser Tyr Ala Cys
                 180   185   190

Arg Phe Thr Glu Thr Ala Arg Leu Arg Gln Glu Arg Ala Ile Gly Arg
                 195   200   205

Pro Val Phe Pro Lys Ala Ile Gly Glu Leu His Gly His Ser Gly Ile
             210   215   220

Ala Arg Leu Ala Leu Phe Tyr Ala Arg Ala Ala Cys Gly Leu Val Asp
225   230   235   240

Pro Ser Ala Gly Thr Gly Val Ala Asn Ala Gly Leu Val Tyr Phe Asn
                 245   250   255

Gly Arg Leu Leu Ala Met Ser Glu Asp Asp Leu Pro Tyr His Val Arg
                 260   265   270

Val Ala Asp Asp Gly Asp Leu Glu Thr Val Gly Arg Tyr Asp Phe Asp
                 275   280   285

Gly Gln Leu Gly Cys Ala Met Ile Ala His Pro Lys Leu Asp Pro Ala
             290   295   300

Thr Gly Glu Leu His Ala Leu Ser Tyr Asp Val Ile Lys Arg Pro Tyr
305   310   315   320

Leu Lys Tyr Phe Tyr Phe Arg Pro Asp Gly Thr Lys Ser Asp Asp Val
                 325   330   335

Glu Ile Pro Leu Glu Gln Pro Thr Met Ile His Asp Phe Ala Ile Thr
             340   345   350

Glu Asn Phe Val Val Val Pro Asp His Gln Val Val Phe Lys Leu Gln
    355   360   365

Glu Met Leu Arg Gly Gly Ser Pro Val Val Leu Asp Lys Glu Lys Thr
    370   375   380

Ser Arg Phe Gly Val Leu Pro Lys His Ala Ala Asp Ala Ser Glu Met
385   390   395   400

Ala Trp Val Asp Val Pro Asp Cys Phe Cys Phe His Leu Trp Asn Ala
         405   410   415

Trp Glu Asp Glu Ala Thr Gly Glu Val Val Val Ile Gly Ser Cys Met
      420   425   430

Thr Pro Ala Asp Ser Ile Phe Asn Glu Ser Asp Glu Arg Leu Glu Ser
      435   440   445

Val Leu Thr Glu Ile Arg Leu Asp Ala Arg Thr Gly Arg Ser Thr Arg
    450   455   460

Arg Ala Val Leu Pro Pro Ser Gln Gln Glu Asn Leu Glu Val Gly Met
465   470   475   480

Val Asn Arg Asn Leu Leu Gly Arg Glu Ser Arg Tyr Ala Tyr Leu Ala
      485   490   495

Val Ala Glu Pro Trp Pro Lys Glu Ser Gly Phe Ala Lys Glu Asp Leu
    500   505   510

Ser Thr Gly Glu Leu Thr Lys Phe Glu Tyr Gly Glu Gly Arg Phe Gly
    515   520   525

Gly Glu Pro Cys Phe Val Pro Met Asp Pro Ala Ala Ala His Pro Arg
    530   535   540

Gly Glu Asp Asp Gly Tyr Val Leu Thr Phe Val His Asp Glu Arg Ala
545   550   555   560

```
Gly Thr Ser Glu Leu Leu Val Val Asn Ala Ala Asp Ile Arg Leu Glu
            565             570             575
```

```
Ala Thr Val Gln Leu Pro Ser Arg Val Pro Phe Gly Phe His Gly Thr
            580             585             590
```

```
Phe Ile Thr Gly Gln Glu Leu Glu Ala Gln Ala Ala
            595             600
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<210> 15
<211> 1818
<212> DNA
<213> Lycopersicon esculentum

<220>
<221> CDS
<222> (1)..(1818)

<400> 15

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Met Ala Thr Thr Thr Ser His Ala Thr Asn Thr Trp Ile Lys Thr Lys
 1               5              10              15
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ttg tca atg cca tca tca aag gag ttt ggt ttt gca tca aac tct att    96
Leu Ser Met Pro Ser Ser Lys Glu Phe Gly Phe Ala Ser Asn Ser Ile
            20              25              30
```

```
tct cta ctc aaa aat caa cat aat agg caa agt ctc aac att aat tcc   144
Ser Leu Leu Lys Asn Gln His Asn Arg Gln Ser Leu Asn Ile Asn Ser
            35              40              45
```

```
tct ctt caa gct cca cct ata ctt cat ttt cct aaa caa tct tca aat   192
Ser Leu Gln Ala Pro Pro Ile Leu His Phe Pro Lys Gln Ser Ser Asn
        50              55              60
```

```
tat caa aca cca aag aat aat aca att tca cac cca aaa caa gaa aac   240
Tyr Gln Thr Pro Lys Asn Asn Thr Ile Ser His Pro Lys Gln Glu Asn
65              70              75              80
```

```
aac aac tcc tct tct tct tca act tcc aag tgg aat tta gtg cag aaa     288
Asn Asn Ser Ser Ser Ser Ser Thr Ser Lys Trp Asn Leu Val Gln Lys
                85              90                  95

gca gca gca atg gct tta gat gct gta gaa agt gct tta act aaa cat     336
Ala Ala Ala Met Ala Leu Asp Ala Val Glu Ser Ala Leu Thr Lys His
            100             105                 110

gaa ctt gaa cac cct ttg ccg aaa aca gcc gac cca cga gtc cag att     384
Glu Leu Glu His Pro Leu Pro Lys Thr Ala Asp Pro Arg Val Gln Ile
            115             120                 125

tct ggg aat ttt gct ccg gta ccg gaa aat cca gtc tgt caa tct ctt     432
Ser Gly Asn Phe Ala Pro Val Pro Glu Asn Pro Val Cys Gln Ser Leu
        130             135             140

ccg gtc acc gga aaa ata ccc aaa tgt gtt caa ggc gtt tac gtt cga     480
Pro Val Thr Gly Lys Ile Pro Lys Cys Val Gln Gly Val Tyr Val Arg
145             150             155                 160

aac gga gct aac cct ctt ttt gaa cca acc gcc gga cac cat ttc ttc     528
Asn Gly Ala Asn Pro Leu Phe Glu Pro Thr Ala Gly His His Phe Phe
                165             170             175

gac ggc gac ggt atg gtt cac gcc gtt caa ttc aaa aat ggg tcg gct     576
Asp Gly Asp Gly Met Val His Ala Val Gln Phe Lys Asn Gly Ser Ala
            180             185             190

agt tac gct tgc cgt ttc act gaa aca gag agg ctt gtt caa gaa aaa     624
Ser Tyr Ala Cys Arg Phe Thr Glu Thr Glu Arg Leu Val Gln Glu Lys
            195             200             205

gct ttg ggt cgc cct gtt ttc cct aaa gcc att ggt gaa tta cat ggt     672
Ala Leu Gly Arg Pro Val Phe Pro Lys Ala Ile Gly Glu Leu His Gly
        210             215             220

cac tct gga att gca agg ctt atg ctg ttt tac gct cgt ggg ctc ttc     720
His Ser Gly Ile Ala Arg Leu Met Leu Phe Tyr Ala Arg Gly Leu Phe
225             230             235             240
```

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gga ctt gtt gat cac agt aaa gga act ggt gtt gca aac gcc ggt tta    768
Gly Leu Val Asp His Ser Lys Gly Thr Gly Val Ala Asn Ala Gly Leu
                245                 250                 255

gtc tat ttc aat aac cga tta ctt gct atg tct gaa gat gat ttg cct    816
Val Tyr Phe Asn Asn Arg Leu Leu Ala Met Ser Glu Asp Asp Leu Pro
                260                 265                 270

tac cat gta aag gta aca ccc acc ggc gat ctt aaa aca gag ggt cga    864
Tyr His Val Lys Val Thr Pro Thr Gly Asp Leu Lys Thr Glu Gly Arg
                275                 280                 285

ttc gat ttc gac ggc cag cta aaa tcc acc atg ata gct cac cca aag    912
Phe Asp Phe Asp Gly Gln Leu Lys Ser Thr Met Ile Ala His Pro Lys
        290                 295                 300

ctc gac cca gtt tcc ggt gag cta ttt gct ctt agc tac gat gtg att    960
Leu Asp Pro Val Ser Gly Glu Leu Phe Ala Leu Ser Tyr Asp Val Ile
305                 310                 315                 320

cag aag cca tac ctc aag tac ttc aga ttt tca aaa aat ggg gaa aaa   1008
Gln Lys Pro Tyr Leu Lys Tyr Phe Arg Phe Ser Lys Asn Gly Glu Lys
                325                 330                 335

tca aat gat gtt gaa att cca gtt gaa gac cca aca atg atg cat gat   1056
Ser Asn Asp Val Glu Ile Pro Val Glu Asp Pro Thr Met Met His Asp
                340                 345                 350

ttc gca att act gag aac ttc gtc gtc att cct gat caa caa gtc gtt   1104
Phe Ala Ile Thr Glu Asn Phe Val Val Ile Pro Asp Gln Gln Val Val
                355                 360                 365

ttc aag atg tct gaa atg atc cgt gga ggt tca ccg gtg gtt tac gac   1152
Phe Lys Met Ser Glu Met Ile Arg Gly Gly Ser Pro Val Val Tyr Asp
        370                 375                 380

aag aac aaa gtt tcc cga ttt ggt att ctg gat aag tac gcg aaa gat   1200
Lys Asn Lys Val Ser Arg Phe Gly Ile Leu Asp Lys Tyr Ala Lys Asp
385                 390                 395                 400
```

ggg tct gat ttg aaa tgg gtt gaa gta cct gat tgt ttc tgt ttc cac   1248
Gly Ser Asp Leu Lys Trp Val Glu Val Pro Asp Cys Phe Cys Phe His
             405   410   415

ctc tgg aat gct tgg gaa gaa gca gaa aca gat gaa atc gtt gta att   1296
Leu Trp Asn Ala Trp Glu Glu Ala Glu Thr Asp Glu Ile Val Val Ile
             420   425   430

ggt tca tgt atg aca cca cca gac tcc att ttc aat gaa tgt gat gaa   1344
Gly Ser Cys Met Thr Pro Pro Asp Ser Ile Phe Asn Glu Cys Asp Glu
             435   440   445

ggg cta aag agt gtt tta tcc gaa atc cgt ctc aat ttg aaa aca ggg   1392
Gly Leu Lys Ser Val Leu Ser Glu Ile Arg Leu Asn Leu Lys Thr Gly
          450   455   460

aaa tca aca aga aaa tcc ata atc gaa aac ccg gat gaa caa gtg aat   1440
Lys Ser Thr Arg Lys Ser Ile Ile Glu Asn Pro Asp Glu Gln Val Asn
465   470   475   480

tta gaa gct gga atg gtg aac cga aac aaa ctc gga agg aaa aca gag   1488
Leu Glu Ala Gly Met Val Asn Arg Asn Lys Leu Gly Arg Lys Thr Glu
             485   490   495

tat gct tat ttg gct atc gct gaa cca tgg cca aaa gtt tct ggt ttt   1536
Tyr Ala Tyr Leu Ala Ile Ala Glu Pro Trp Pro Lys Val Ser Gly Phe
             500   505   510

gca aaa gta aac ctg ttc acc ggt gaa gtt gag aaa ttc att tat ggt   1584
Ala Lys Val Asn Leu Phe Thr Gly Glu Val Glu Lys Phe Ile Tyr Gly
          515   520   525

gac aac aaa tat ggt ggg gaa cct ctt ttt tta cca aga gac ccc aac   1632
Asp Asn Lys Tyr Gly Gly Glu Pro Leu Phe Leu Pro Arg Asp Pro Asn
          530   535   540

agc aag gaa gaa gac gat ggt tat att tta gct ttc gtt cac gat gag   1680
Ser Lys Glu Glu Asp Asp Gly Tyr Ile Leu Ala Phe Val His Asp Glu
545   550   555   560

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aaa gaa tgg aaa tca gaa ctg caa att gtt aac gca atg agt ttg aag    1728
Lys Glu Trp Lys Ser Glu Leu Gln Ile Val Asn Ala Met Ser Leu Lys
                565                 570                 575


ttg gag gca act gtg aag ctt cca tca aga gtt cct tat gga ttt cat    1776
Leu Glu Ala Thr Val Lys Leu Pro Ser Arg Val Pro Tyr Gly Phe His
                580                 585                 590


gga aca ttc ata aac gcc aat gat ttg gca aat cag gca tga           1818
Gly Thr Phe Ile Asn Ala Asn Asp Leu Ala Asn Gln Ala
                595                 600                 605



<210> 16
<211> 605
<212> PRT
<213> Lycopersicon esculentum

<400> 16
Met Ala Thr Thr Thr Ser His Ala Thr Asn Thr Trp Ile Lys Thr Lys
  1                 5                  10                  15


Leu Ser Met Pro Ser Ser Lys Glu Phe Gly Phe Ala Ser Asn Ser Ile
                20                  25                  30


Ser Leu Leu Lys Asn Gln His Asn Arg Gln Ser Leu Asn Ile Asn Ser
               35                  40                  45


Ser Leu Gln Ala Pro Pro Ile Leu His Phe Pro Lys Gln Ser Ser Asn
            50                  55                  60


Tyr Gln Thr Pro Lys Asn Asn Thr Ile Ser His Pro Lys Gln Glu Asn
 65                 70                  75                  80


Asn Asn Ser Ser Ser Ser Ser Thr Ser Lys Trp Asn Leu Val Gln Lys
                85                  90                  95


Ala Ala Ala Met Ala Leu Asp Ala Val Glu Ser Ala Leu Thr Lys His
                100                 105                 110
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Glu Leu Glu His Pro Leu Pro Lys Thr Ala Asp Pro Arg Val Gln Ile
  115   120   125

Ser Gly Asn Phe Ala Pro Val Pro Glu Asn Pro Val Cys Gln Ser Leu
  130   135   140

Pro Val Thr Gly Lys Ile Pro Lys Cys Val Gln Gly Val Tyr Val Arg
145   150   155   160

Asn Gly Ala Asn Pro Leu Phe Glu Pro Thr Ala Gly His His Phe Phe
     165   170   175

Asp Gly Asp Gly Met Val His Ala Val Gln Phe Lys Asn Gly Ser Ala
     180   185   190

Ser Tyr Ala Cys Arg Phe Thr Glu Thr Glu Arg Leu Val Gln Glu Lys
  195   200   205

Ala Leu Gly Arg Pro Val Phe Pro Lys Ala Ile Gly Glu Leu His Gly
  210   215   220

His Ser Gly Ile Ala Arg Leu Met Leu Phe Tyr Ala Arg Gly Leu Phe
225   230   235   240

Gly Leu Val Asp His Ser Lys Gly Thr Gly Val Ala Asn Ala Gly Leu
     245   250   255

Val Tyr Phe Asn Asn Arg Leu Leu Ala Met Ser Glu Asp Asp Leu Pro
     260   265   270

Tyr His Val Lys Val Thr Pro Thr Gly Asp Leu Lys Thr Glu Gly Arg
  275   280   285

Phe Asp Phe Asp Gly Gln Leu Lys Ser Thr Met Ile Ala His Pro Lys
  290   295   300

Leu Asp Pro Val Ser Gly Glu Leu Phe Ala Leu Ser Tyr Asp Val Ile
305   310   315   320

Gln Lys Pro Tyr Leu Lys Tyr Phe Arg Phe Ser Lys Asn Gly Glu Lys

                325   330   335

Ser Asn Asp Val Glu Ile Pro Val Glu Asp Pro Thr Met Met His Asp
          340   345   350

Phe Ala Ile Thr Glu Asn Phe Val Val Ile Pro Asp Gln Gln Val Val
          355   360   365

Phe Lys Met Ser Glu Met Ile Arg Gly Gly Ser Pro Val Val Tyr Asp
      370   375   380

Lys Asn Lys Val Ser Arg Phe Gly Ile Leu Asp Lys Tyr Ala Lys Asp
385   390   395   400

Gly Ser Asp Leu Lys Trp Val Glu Val Pro Asp Cys Phe Cys Phe His
              405   410   415

Leu Trp Asn Ala Trp Glu Glu Ala Glu Thr Asp Glu Ile Val Val Ile
          420   425   430

Gly Ser Cys Met Thr Pro Pro Asp Ser Ile Phe Asn Glu Cys Asp Glu
          435   440   445

Gly Leu Lys Ser Val Leu Ser Glu Ile Arg Leu Asn Leu Lys Thr Gly
      450   455   460

Lys Ser Thr Arg Lys Ser Ile Ile Glu Asn Pro Asp Glu Gln Val Asn
465   470   475   480

Leu Glu Ala Gly Met Val Asn Arg Asn Lys Leu Gly Arg Lys Thr Glu
              485   490   495

Tyr Ala Tyr Leu Ala Ile Ala Glu Pro Trp Pro Lys Val Ser Gly Phe
              500   505   510

Ala Lys Val Asn Leu Phe Thr Gly Glu Val Glu Lys Phe Ile Tyr Gly
      515   520   525

Asp Asn Lys Tyr Gly Gly Glu Pro Leu Phe Leu Pro Arg Asp Pro Asn
      530   535   540

Ser Lys Glu Glu Asp Asp Gly Tyr Ile Leu Ala Phe Val His Asp Glu
545   550   555   560

Lys Glu Trp Lys Ser Glu Leu Gln Ile Val Asn Ala Met Ser Leu Lys
                 565   570   575

Leu Glu Ala Thr Val Lys Leu Pro Ser Arg Val Pro Tyr Gly Phe His
             580   585   590

Gly Thr Phe Ile Asn Ala Asn Asp Leu Ala Asn Gln Ala
             595   600   605

<210> 17
<211> 1617
<212> DNA
<213> Arabidopsis thaliana

<220>
<221> CDS
<222> (1)..(1617)

<400> 17
atg gcg gag aaa ctc agt gat ggc agc atc atc atc tca gtc cat cct   48
Met Ala Glu Lys Leu Ser Asp Gly Ser Ile Ile Ile Ser Val His Pro
 1   5   10   15

aga ccc tcc aag ggt ttc tcc tcg aag ctt ctc gat ctt ctc gag aga   96
Arg Pro Ser Lys Gly Phe Ser Ser Lys Leu Leu Asp Leu Leu Glu Arg
            20   25   30

ctt gtc gtc aag ctc atg cac gat gct tct ctc cct ctc cac tac ctc   144
Leu Val Val Lys Leu Met His Asp Ala Ser Leu Pro Leu His Tyr Leu
            35   40   45

tca ggc aac ttc gct ccc atc cgt gat gaa act cct ccc gtc aag gat   192
Ser Gly Asn Phe Ala Pro Ile Arg Asp Glu Thr Pro Pro Val Lys Asp
    50   55   60

ctc ccc gtc cat gga ttt ctt ccc gaa tgc ttg aat ggt gaa ttt gtg   240
Leu Pro Val His Gly Phe Leu Pro Glu Cys Leu Asn Gly Glu Phe Val
65   70   75   80

agg gtt ggt cca aac ccc aag ttt gat gct gtc gct gga tat cac tgg   288
Arg Val Gly Pro Asn Pro Lys Phe Asp Ala Val Ala Gly Tyr His Trp
              85   90   95

ttt gat gga gat ggg atg att cat ggg gta cgc atc aaa gat ggg aaa   336
Phe Asp Gly Asp Gly Met Ile His Gly Val Arg Ile Lys Asp Gly Lys
              100   105   110

gct act tat gtt tct cga tat gtt aag aca tca cgt ctt aag cag gaa   384
Ala Thr Tyr Val Ser Arg Tyr Val Lys Thr Ser Arg Leu Lys Gln Glu
            115   120   125

gag ttc ttc gga gct gcc aaa ttc atg aag att ggt gac ctt aag ggg   432
Glu Phe Phe Gly Ala Ala Lys Phe Met Lys Ile Gly Asp Leu Lys Gly
           130   135   140

ttt ttc gga ttg cta atg gtc aat atc caa cag ctg aga acg aag ctc   480
Phe Phe Gly Leu Leu Met Val Asn Ile Gln Gln Leu Arg Thr Lys Leu
145   150   155   160

aaa ata ttg gac aac act tat gga aat gga act gcc aat aca gca ctc   528
Lys Ile Leu Asp Asn Thr Tyr Gly Asn Gly Thr Ala Asn Thr Ala Leu
              165   170   175

gta tat cac cat gga aaa ctt cta gca tta cag gag gca gat aag ccg   576
Val Tyr His His Gly Lys Leu Leu Ala Leu Gln Glu Ala Asp Lys Pro
              180   185   190

tac gtc atc aaa gtt ttg gaa gat gga gac ctg caa act ctt ggt ata   624
Tyr Val Ile Lys Val Leu Glu Asp Gly Asp Leu Gln Thr Leu Gly Ile
           195   200   205

ata gat tat gac aag aga ttg acc cac tcc ttc act gct cac cca aaa   672
Ile Asp Tyr Asp Lys Arg Leu Thr His Ser Phe Thr Ala His Pro Lys
           210   215   220

gtt gac ccg gtt acg ggt gaa atg ttt aca ttc ggc tat tcg cat acg   720
Val Asp Pro Val Thr Gly Glu Met Phe Thr Phe Gly Tyr Ser His Thr
225   230   235   240

cca cct tat ctc aca tac aga gtt atc tcg aaa gat ggc att atg cat   768
Pro Pro Tyr Leu Thr Tyr Arg Val Ile Ser Lys Asp Gly Ile Met His
           245   250   255

gac cca gtc cca att act ata tca gag cct atc atg atg cat gat ttt   816
Asp Pro Val Pro Ile Thr Ile Ser Glu Pro Ile Met Met His Asp Phe
           260   265   270

gct att act gag act tat gca atc ttc atg gat ctt cct atg cac ttc   864
Ala Ile Thr Glu Thr Tyr Ala Ile Phe Met Asp Leu Pro Met His Phe
           275   280   285

agg cca aag gaa atg gtg aaa gag aag aaa atg ata tac tca ttt gat   912
Arg Pro Lys Glu Met Val Lys Glu Lys Lys Met Ile Tyr Ser Phe Asp
  290   295   300

ccc aca aaa aag gct cgt ttt ggt gtt ctt ccg cgc tat gcc aag gat   960
Pro Thr Lys Lys Ala Arg Phe Gly Val Leu Pro Arg Tyr Ala Lys Asp
305   310   315   320

gaa ctt atg att aga tgg ttt gag ctt ccc aac tgc ttt att ttc cac 1008
Glu Leu Met Ile Arg Trp Phe Glu Leu Pro Asn Cys Phe Ile Phe His
           325   330   335

aac gcc aat gct tgg gaa gaa gag gat gaa gtc gtc ctc atc act tgt 1056
Asn Ala Asn Ala Trp Glu Glu Glu Asp Glu Val Val Leu Ile Thr Cys
          340   345   350

cgt ctt gag aat cca gat ctt gac atg gtc agt ggg aaa gtg aaa gaa 1104
Arg Leu Glu Asn Pro Asp Leu Asp Met Val Ser Gly Lys Val Lys Glu
       355   360   365

aaa ctc gaa aat ttt ggc aac gaa ctg tac gaa atg aga ttc aac atg 1152
Lys Leu Glu Asn Phe Gly Asn Glu Leu Tyr Glu Met Arg Phe Asn Met
     370   375   380

aaa acg ggc tca gct tct caa aaa aaa cta tcc gca tct gcg gtt gat   1200
Lys Thr Gly Ser Ala Ser Gln Lys Lys Leu Ser Ala Ser Ala Val Asp
385   390   395   400

ttc ccc aga atc aat gag tgc tac acc gga aag aaa cag aga tac gta   1248
Phe Pro Arg Ile Asn Glu Cys Tyr Thr Gly Lys Lys Gln Arg Tyr Val
                405   410   415

tat gga aca att ctg gac agt atc gca aag gtt acc gga atc atc aag   1296
Tyr Gly Thr Ile Leu Asp Ser Ile Ala Lys Val Thr Gly Ile Ile Lys
                420   425   430

ttt gat ctg cat gca gaa gct gag aca ggg aaa aga atg ctg gaa gta   1344
Phe Asp Leu His Ala Glu Ala Glu Thr Gly Lys Arg Met Leu Glu Val
            435   440   445

gga ggt aat atc aaa gga ata tat gac ctg gga gaa ggc aga tat ggt   1392
Gly Gly Asn Ile Lys Gly Ile Tyr Asp Leu Gly Glu Gly Arg Tyr Gly
        450   455   460

tca gag gct atc tat gtt ccg cgt gag aca gca gaa gaa gac gac ggt   1440
Ser Glu Ala Ile Tyr Val Pro Arg Glu Thr Ala Glu Glu Asp Asp Gly
465   470   475   480

tac ttg ata ttc ttt gtt cat gat gaa aac aca ggg aaa tca tgc gtg   1488
Tyr Leu Ile Phe Phe Val His Asp Glu Asn Thr Gly Lys Ser Cys Val
                485   490   495

act gtg ata gac gca aaa aca atg tcg gct gaa ccg gtg gca gtg gtg   1536
Thr Val Ile Asp Ala Lys Thr Met Ser Ala Glu Pro Val Ala Val Val
                500   505   510

gag ctg ccg cac agg gtc cca tat ggc ttc cat gcc ttg ttt gtt aca   1584
Glu Leu Pro His Arg Val Pro Tyr Gly Phe His Ala Leu Phe Val Thr
            515   520   525

gag gaa caa ctc cag gaa caa act ctt ata taa   1617
Glu Glu Gln Leu Gln Glu Gln Thr Leu Ile
            530   535

<210> 18
<211> 538
<212> PRT
<213> Arabidopsis thaliana

<400> 18
Met Ala Glu Lys Leu Ser Asp Gly Ser Ile Ile Ile Ser Val His Pro
1   5   10   15

Arg Pro Ser Lys Gly Phe Ser Ser Lys Leu Leu Asp Leu Leu Glu Arg
            20   25   30

Leu Val Val Lys Leu Met His Asp Ala Ser Leu Pro Leu His Tyr Leu
        35   40   45

Ser Gly Asn Phe Ala Pro Ile Arg Asp Glu Thr Pro Pro Val Lys Asp
    50   55   60

Leu Pro Val His Gly Phe Leu Pro Glu Cys Leu Asn Gly Glu Phe Val
65   70   75   80

Arg Val Gly Pro Asn Pro Lys Phe Asp Ala Val Ala Gly Tyr His Trp
            85   90   95

Phe Asp Gly Asp Gly Met Ile His Gly Val Arg Ile Lys Asp Gly Lys
            100   105   110

Ala Thr Tyr Val Ser Arg Tyr Val Lys Thr Ser Arg Leu Lys Gln Glu
        115   120   125

Glu Phe Phe Gly Ala Ala Lys Phe Met Lys Ile Gly Asp Leu Lys Gly
    130   135   140

Phe Phe Gly Leu Leu Met Val Asn Ile Gln Gln Leu Arg Thr Lys Leu
145   150   155   160

Lys Ile Leu Asp Asn Thr Tyr Gly Asn Gly Thr Ala Asn Thr Ala Leu
                165   170   175

Val Tyr His His Gly Lys Leu Leu Ala Leu Gln Glu Ala Asp Lys Pro
        180   185   190

Tyr Val Ile Lys Val Leu Glu Asp Gly Asp Leu Gln Thr Leu Gly Ile
     195   200   205

Ile Asp Tyr Asp Lys Arg Leu Thr His Ser Phe Thr Ala His Pro Lys
  210   215   220

Val Asp Pro Val Thr Gly Glu Met Phe Thr Phe Gly Tyr Ser His Thr
225   230   235   240

Pro Pro Tyr Leu Thr Tyr Arg Val Ile Ser Lys Asp Gly Ile Met His
        245   250   255

Asp Pro Val Pro Ile Thr Ile Ser Glu Pro Ile Met Met His Asp Phe
     260   265   270

Ala Ile Thr Glu Thr Tyr Ala Ile Phe Met Asp Leu Pro Met His Phe
     275   280   285

Arg Pro Lys Glu Met Val Lys Glu Lys Lys Met Ile Tyr Ser Phe Asp
   290   295   300

Pro Thr Lys Lys Ala Arg Phe Gly Val Leu Pro Arg Tyr Ala Lys Asp
305   310   315   320

Glu Leu Met Ile Arg Trp Phe Glu Leu Pro Asn Cys Phe Ile Phe His
        325   330   335

Asn Ala Asn Ala Trp Glu Glu Glu Asp Glu Val Val Leu Ile Thr Cys
     340   345   350

Arg Leu Glu Asn Pro Asp Leu Asp Met Val Ser Gly Lys Val Lys Glu
     355   360   365

Lys Leu Glu Asn Phe Gly Asn Glu Leu Tyr Glu Met Arg Phe Asn Met
     370   375   380

Lys Thr Gly Ser Ala Ser Gln Lys Lys Leu Ser Ala Ser Ala Val Asp

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        385                390                395                400

Phe Pro Arg Ile Asn Glu Cys Tyr Thr Gly Lys Lys Gln Arg Tyr Val
                405                410                415

Tyr Gly Thr Ile Leu Asp Ser Ile Ala Lys Val Thr Gly Ile Ile Lys
                420                425                430

Phe Asp Leu His Ala Glu Ala Glu Thr Gly Lys Arg Met Leu Glu Val
                435                440                445

Gly Gly Asn Ile Lys Gly Ile Tyr Asp Leu Gly Glu Gly Arg Tyr Gly
        450                455                460

Ser Glu Ala Ile Tyr Val Pro Arg Glu Thr Ala Glu Glu Asp Asp Gly
465                470                475                480

Tyr Leu Ile Phe Phe Val His Asp Glu Asn Thr Gly Lys Ser Cys Val
                485                490                495

Thr Val Ile Asp Ala Lys Thr Met Ser Ala Glu Pro Val Ala Val Val
                500                505                510

Glu Leu Pro His Arg Val Pro Tyr Gly Phe His Ala Leu Phe Val Thr
        515                520                525

Glu Glu Gln Leu Gln Glu Gln Thr Leu Ile
        530                535
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<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: artificially
     synthesized sequence

<400> 19

attgaattca tgccttcagc ttcaaac   27

<210> 20
<211> 31
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: artificially
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<400> 20
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<210> 21
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<212> DNA
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<223> Description of Artificial Sequence: artificially
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<400> 21
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<210> 22
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<220>
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<400> 22
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<210> 23
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<210> 24
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<400> 24
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<400> 25
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<210> 26
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<400> 26
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<210> 27
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<400> 27
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<210> 28
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      synthesized sequence

<400> 28
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<210> 29
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      synthesized sequence

<400> 29
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<210> 30
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<210> 31
<211> 32
<212> DNA
<213> Artificial Sequence

<220>
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      synthesized sequence

<400> 31
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<210> 32

<211> 17
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      synthesized sequence

<400> 32
gttttcccag tcacgac   17